

GCTCAGGTTAGCCCGAAAGATGTTTCAGTATAGATTAGGCCACTCTAATTTAATGATCACTGAAAATACTTACTGGCATACTAA  
CCAAGAGAATGCAAAAAAGCCGTCTCAAATTATGAAACAGCTATCAACAATTTATAA

(SEQ. ID. NO. 195)

MEGLVRVHLLPVFGDYKLSKLTTPILQQQVKNKWADKANKGEKGAFANYSLHNMNKRILKYGVAIQVIQYNPANDVIVPRKQQ  
KEKAADVYLDNKELKQFLDYLDALDQSNYENLFDVVLYKTLATGCRISEALALEWSDIDLESGLVISINKTLNRYQEINSPKS  
SAGYRDIPIDKATLLLLKQYKNRQQIQSWKLGRSETVVFTEKYAYACNLRLKRLNKHFDAAGVTNVSFHGFRHTHTTMMLY  
AQVSPKDVQYRLGHSNLMITENTYWHNTQENAKKAVSNYETAINN LZ--

Please delete the first paragraph located on page 101, line 1, and replace therewith the following replacement paragraph. A marked-up version showing changes made to the replacement paragraphs is contained herewith as Appendix B.

--What is claimed is:--

### REMARKS

Claims 1-20 are currently pending in this application. Upon entry of the amendments contained herein, claims 1-22 will be pending. No new matter is introduced into the specification by way of these amendments. The Specification amendments are introduced to conform the application to the requirements of 37 C.F.R. § § 1.77 and 1.821-1.825. Support for the claim amendments can be found, inter alia, throughout the application as originally filed, for example, in the originally filed claims; on page 9, last two full paragraphs; and in Table 3. Support for new claim 21 can be found, inter alia, throughout the specification, for example, on page 9, last two full paragraphs. Support for new claim 22 can be found, inter alia, throughout the Specification, for example, on page 9, last two full paragraphs and on page 13, third full paragraph.

### Priority

The Examiner has made acknowledgement of Applicant's claim of foreign priority based on "German" foreign application number 9816337.1. Applicants respectfully state that the claim to foreign priority is in fact based upon Great Britain priority application number 9816337.1. Applicants respectfully direct the Examiner's attention to the executed declaration, claiming the benefit of GB 9816337.1, as well as the cover sheet of International Publication Number WO 00/06738 (reciting the priority application GB 9816337.1, and of which the instant application claims benefit), attached herewith as Appendix C. Applicants will provide the Examiner with a certified copy of Great Britain priority application GB 9816337.1 within a reasonable time of receipt of the document.

### Restriction Requirement

In the Office Action dated September 22, 2002, the Examiner required restriction of the pending claims under 35 U.S.C. § 121 between one of the following inventions:

- “1.) Claims 7-8 and 13 drawn to DNA, host cell and a vaccine composition, classified in class 536, subclass 23.7. Further election of invention required.
- 2.) Claims 1-6 and 10-12 drawn to polypeptide, classified in class 424, subclass 190.1. Further election of invention required.
- 3.) Claims 15-16 drawn to an antibody classified in class 530, subclass 388.6. Further election of invention required.
- 4.) Claim 9 drawn to a method for inducing immune response using polypeptide classified in class 424, subclass 165.1. Further election of invention required.
- 5.) Claim 20 drawn to a method of use for the treatment or prophylaxis of *S. pneumoniae* infection using protein or polypeptide classified in class 424, subclass 184.1. Further election of invention required.
- 6.) Claim 19 drawn to a method of determining whether a protein is anti-microbial target in vivo or in vitro classified in class 424, subclass 93.1. Further election of invention required.
- 7.) Claim 18 drawn to a method of detecting *S. pneumoniae* infection using nucleic acid, SEQ ID NO:1 classified in class 435, subclass 6. Further election of invention required.
- 8.) Claim 14 drawn to a method of detecting *S. pneumoniae* infection using protein antigen, SEQ ID NO:2, classified in class 435, subclass 7.22. Further election of invention required.
- 9.) Claim 17 drawn to a method of detecting *S. pneumoniae* infection using antibody classified in class 435, subclass 7.22. Further election of invention required.”

See Paper No. 11, page 2. The Examiner further required election of one sequence falling within one of the above-identified inventions.

Applicants were contacted by the Examiner and invited to make an election to a restriction requirement by telephone. The Examiner originally designated (by telephone) as separate inventions the following 5 groups: (I) *S. pneumoniae* nucleic acid sequences (claims 7-8 and 18); (II) *S. pneumoniae* peptide sequences (claims 1-6 and 9-14); (III) antibodies to *S. pneumoniae* antigens (claims 15-17); (IV) methods of assay design (claim 19); and therapeutic methods (claim 20). Based upon this grouping of claims, Applicants provisionally elected by telephone on July 09, 2002, without traverse, the invention of Group V, claim 20, *drawn to therapeutic methods*. To the extent that the restriction requirement in the Office Action dated September 20, 2002, differs from that described in the telephone restriction to which Applicants responded on July 9, 2002, the restriction is traversed. In particular, Applicants submit that there is substantial overlap in scope between Group 4 and Group 5 or Group 6. Dependent claims to particular embodiments may properly fall within the scope of both generic claim 9 and generic claim 20. Likewise, dependent claims to other

embodiments may properly fall within the scope of both generic claim 9 and generic claim 19. Applicants respectfully request that the restriction requirement be recast.

Applicants also submit that the requirement for election of a single nucleotide (polypeptide) sequence is properly characterized as an election of species. Applicants provisionally elect the invention of claim 20, now classified in Group 5.

Consistent with the requirement of election of species, Applicants hereby further elect the protein sequence as set forth in SEQ ID NO:162.

### **Rejections**

#### *35 U.S.C. § 101*

The Examiner rejected claim 20 under 35 U.S.C. § 101, as allegedly improperly defining a process. Applicants have amended claim 20 to conform with U.S. Rules of Practice, thereby rendering the Examiner's rejection moot. Accordingly, Applicants respectfully request reconsideration and withdrawal of the rejection of claim 20 under 35 U.S.C. § 101.

#### *35 U.S.C. § 112, 2<sup>nd</sup> Paragraph*

The Examiner rejected claim 20 under 35 U.S.C. § 112, 2<sup>nd</sup> Paragraph as allegedly indefinite for failing to particularly point out and distinctly claim the subject matter which Applicants regard as the invention. More specifically, the Examiner stated that "the claim does not set forth any steps involved in the method/process, [and therefore] it is unclear what method/process applicant is intending to encompass." *See* Paper No. 11, Page 7, lines 4-6. Similarly, the Examiner stated that the claim is further indefinite because "the claims do not recite positive active steps so that the claims will set out and circumscribe particular area with reasonable degree of precision and particularly [sic] and make clear what subject matter the claims encompass, as well as make clear the subject matter from which others would be precluded." *See* Paper No. 11, Page 7, lines 9-13.

The Examiner further determined that "additional structural parameters, such as amino acid sequence of the polypeptide (for example; SEQ.ID.NO), would enable more definitive identification of the claimed polypeptide without ambiguity." *See* Paper No. 11, Page 7, lines 17-19.

The Examiner also rejected claim 20 as allegedly vague for the recitation of "capable of" within the claim, as this recitation allegedly makes it difficult to determine the metes and bounds of the claim.

Applicants have amended claim 20 to confirm to U.S. practice. As a result of the claim amendment, the Examiner's rejection of claim 20 has been rendered moot. Applicants respectfully request withdrawal of the rejection of claim 20 under 35 U.S.C. § 112, 2<sup>nd</sup> Paragraph, and reconsideration of the application in view of the amendments.

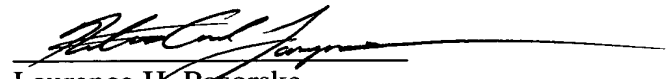
**Conclusion**

Applicants believe that amendment of the subject application has placed this application in a condition for allowance. Early notification of a favorable consideration is respectfully requested.

Respectfully submitted,

Dated: March 20, 2003

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**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE**

In re Patent Application of :

GILBERT, C.F.G., *et al.*

Serial No.: 09/769,787

Filed: January 26, 2001

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Examiner: Baskar, P.

Group Art Unit: 1645

For: NUCLEIC ACIDS AND PROTEINS FROM STREPTOCOCCUS PNEUMONIAE

Under Secretary of Commerce for Intellectual Property and  
Director of the United States Patent and Trademark Office

Washington, D.C. 20231

**Appendix A - Version with Markings to show changes made to the Claims**

In claim 20:

--20. (Amended) A method for treatment or prophylaxis of *S. pneumoniae* infection comprising administering to a patient in need thereof ~~The use of~~ an agent capable of antagonizing, inhibiting or otherwise interfering with the function or expression of a protein or polypeptide as defined in Tables 2-4 ~~in the manufacture of a medicament for use in the treatment or prophylaxis of *S. pneumoniae* infection,~~ wherein said agent induces an immune response in said patient to a polypeptide having a sequence comprising SEQ. ID NO:162.--

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**Appendix B - Version with Markings to show changes made**

In the paragraphs starting at page 29, 1<sup>st</sup> full paragraph (immediately below the heading  
"Table 1") and extending to and including the last full paragraph located on page 100:

**4101.1 (SEQ. ID. NO. 208)**

ATGGAAGAGTTAGTGACCTTAGATTGTTTGTATTGACAGAACTAAGATTGAAGCCAATGCCAACAAGTATAGTTTTGTGTG  
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CCTGCTTTGGGCAGATAAAGGCTTCTTTGGGTTACAAGAGATGTAATCTGAGAGGGAAGCGTCAAGTGAGAATTGACATGG  
GATTGGTACTTATGGCCAATAACCTCCTAAAATATAGTAAAATGAAATAA

**4101.3 (SEQ. ID. NO. 209)**

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GTCTGCTAAACCGGAAGAAAAAGCTCCAAGGATAGAATCACAAGCTTCAAAATCAAGAAAAACCGCTCAAGGAAGATGCTAA  
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GTCAGTTAGTGGGGCATTATCCAAATGGTTATAACCAAGTTCTCATATGATATCACCAAATACCTTCAAAAAGATGGTGGTGA  
GAATGTGATTGCTGTCCATGCAGTCAACAAACAGCCAAGTAGCCGTTGGTATTTCAGGAAGTGGTATCTATCGTGATGTGACT  
TTACAAGTGACAGATAAGGTGCATGTTGAGAAAAATGGGACAACTATTTTAACACCAAACTTGAAGAACAAACAACATGGC  
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AACGAGGTGGTCATGCTGTAAACAGGCTTAGTTCGTACAGCGAGTCGTACCTTAAAGACATGAATCAACAAGCCTAGATGC  
GATTTTGAAGATTGAAGACCAAACTCTGGACTGTTTTAAATGACAAAACCTGCCTTGTACGAATTGATTACGCGTGTITACC  
GTGACGGTCAATTGGTTGATGCTAAGAAGGATTTGTTTGGTTACCGTTACTATCACTGGACTCCAAATGAAGGTTTCTCTTTG  
AATGGTGAACTGATTAATTTCCATGGAGTATCCTTGCACCACGACCATGGGGCGCTTGGAGCAGAAGAAAACCTATAAAGCA  
GAATATCGCCGCTCAAAACAAATGAAGGAGATGGGAGTTAACTCCATCCGTACAACCCACAACCTGCTAGTGAGCAAACT  
TGCAAAATCGCAGCAGAACTAGGTTTACTCGTTACGGAAGAGGCCCTTTGATACGTGGTATGGTGGCAAGAAACCTTATGACTA  
TGGACGTTTCTTTGAAAAAGATGCCACTACCCAGAAGCTCGAAAAGGTGAAAAATGGTCTGATTTTGACCTACGTACCATG



GTCTGAAAGAGGCCAAAAACAACCCCTGCTATCTTCATGTGGTCAATTGGTAATGAAATAGGTGAAGCTAATGGTGATGCCCACT  
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4101.5 (SEQ. ID. NO. 210)

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## 4102.1 (SEQ. ID. NO. 211)

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## 4106.1 (SEQ. ID. NO. 212)

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 AGTTTACTACTATTACCAAGTTATCATCGCAAAACAACGACTATCTGAAAGATAACAAAGAAGAAGCTCGCAAGTCAATCCA  
 AGCCATCAAAAAAGGCTACCAATATGCCATGGAACATCCAGAAGAAGCTGCAGATATTCTCATCAAGAATGCACCTGAACTC  
 AAGGAAAAACGTGACTTTGTCATCGAATCTCAAAAAATCTGTCAAAAGAATACGCAAGCGACAAGGAAAAAATGGGGTCAA  
 TTTGACGCAAGCTCGCTGGAATGCTTTCTACAAATGGGATAAAGAAAAATGGTATCCTTAAAGAAGACTTGACAGACAAAGGCT  
 TCACCAACGAATTTGTGAAATAA

## 4106.4 (SEQ. ID. NO. 213)

ATGATAAAAAATCCTAAATTATTAACCAAGTCTTTTTTAAGAAGTTTTGCAATTTCTAGGTGGTGTGGTCTAGTCATTCATAT  
 AGCTATTTATTTGACCTTTCTTTTTATTATATTCAACTGGAGGGGAAAAAGTTAATGAGAGCGCAAGAGTGTTTACGGAGT  
 ATTTAAAGACTAAGACATCTGATGAAATTTCAAGCTTACTCCAGTCTTATTCAAAGTCTTGACCATATCTGCTCACCTTAAAA  
 AGAGATATTGTAGATAAGCGGCTCCCTCTTGTGCATGACTTGGATATTAAGATGGAAAGCTATCAAATATATCGTGATGTT  
 AGATATGCTGTTAGTACAGCAGATGGTAAACAGGTAACCGTGCAATTTGTTACGGGGTGGATGTCTACAAAGAAGCAAG  
 AATATTTGCTTTTGTATCTTTCCATATACATTTTGGTTACAAATGCTTTTCTTTGTTTTTCTTTTCTTATTTTATACTAAAGCTTT  
 GCTCAATCCTCTTTTTTACATTTTCAAGAGTACTAGTAAATGCAAGATTTGGATGACAATATTCGTTTTGATGAAAGTAGGA  
 AAGATGAAGTTGGTGAAGTTGGAAAAACAGATTAATGGTATGTATGAGCACTTGTGAAGGTTATTTATGAGTTGGAAAGTCTG  
 TAATGAGCAAAATGTAATAATGCAAAATCAAAAAGGTTTCTTTGTCGCGGAGCATCACATGAGTTGAAAAACCCCTTTAGCC  
 AGTCTTAGAATTAATCCTAGAGAATATGCAGCATAATATTGGAGATTACAAAGATCATCCAAAATATATTGCAAAAGAGTATAA  
 ATAAGATTGACCAGATGAGCCACTTATTAGAAGAAGTACTGGAGTCTTCTAAATCCAAGAGTGGACAGAGTGTCTGTGAGAC  
 CTTGACTGTTAAGCCAGTTTATGATGATATTTATCACGTTATCAAGAATTAGCTCAATTAAGGTGTTACAATTGAAAAATC  
 AATTGACAGATGCTACAGGGTCTCATGAGTCTTAGGCGATTGATAAGGTTTGGACAAACCTGATTAGTAATGCAATTA  
 ATATTCAGATAAAAAATGGGCGTGAATCATATCCGAGCAAGATGGCTATCTCTCTATCAAAAAATACATGTGCGCCTCTAAGT  
 GACCAAGAAGTACAACATTTATTGATATATTCTATCATTTCTCAAATCGTGACAGATAAGGATGAAAGTTCCGGTTTGGGTCT  
 TTACATTGTGAATAATATTTTAGAAAGCTATCAAATGGATTATAGTTTTCTCCCTTATGAACACGGTATGGAATTTAAGATTA  
 GCTGTAG

## 4106.6 (SEQ. ID. NO. 214)

ATGTATTAGGAGATTGATGGAGAAAGCCGAGTGTGGTCAATTTTCAATACTTTCTTTCTATTACAAGAGTCTCAGACGAC  
 CGTCAAGGCTGTAATGGAAGAAACAGGATTTTCAAAGCAACCCTAACCAAAATATGTCACCCTGCTCAATGACAAGGCTTTG  
 GATAGTGGCTTAGAGCTGGCTATTCACCTCAGAAGATGAAAACTGCGTCTGTCTATCGGTGCAGCTACCAAGGGGAGAGATA  
 TTCGGAGCTTGTGTTTGGAGAGTGTGTTAAATACCAAGATTTTGGTTATCTTTCTCTACCAACCAACAGTTTTAGCCCATCAGC  
 TGGCTCAAGAATTGGTGATTAGCGAGGCTACGCTTGGTCTGCTGCTGTTTAAATCAGATTTTTGTCAAGATTTGATTATA  
 TCCATCAAAAATGGCCGTTGGCGAGGTCCAGAGCATCAGATTCACTATTTCTATTTCTGCTTTTCCGAAAGGTCTGGTTCGAG  
 TCAGGAATTGGGAAGGTACATGCAGAAACCAAGAGAGAAAAACAGGAGATTGCCAATTTAGAGGAAATCTGCGGTGCAAGTTT  
 GTCTGCGGGGCAAAAATGGACTTGGTTCTCTGGGCTCACATCAGTCAACAACAGTCTTCCGGTCAATGCTTCAGTTTCAAG  
 TCATAGAAAGCAAAAATGCGAGGGTATTTGACAATCTTTTATCTTCGTTTGGTGAGAAAGGTTCCGTCCTTTTGTCTGGG  
 CAACATATTCACCTAGGAGTTGAGGATGGTGAGATGATGATATCTTCTCTTTCTCTCTATCTCATCGCATTCTCTCTTCAT



ACTATGGAGTATATTCTTGGTTTTGGAGGGCAGTTGGCAGATTTACTGACGCAATTGATTCAAGAAATGAAGAAGGAGGAAC  
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TTACAGGATCGCTACAAGTACCAGTTAGAGAATCGTCATCCATATTTACTGATGGAACATGATTTTAAAGAGACAGCAGAGG  
AGATTTTTCATGCTCTACCTGCTTTTCAACAGGGGACAGATTTAGATAAGAAGATTCTCTGGGAATGGCTCCAGTTAATCGAA  
TATATGGCTGAAAACGGTGGCCAGCATATGCGGATTGGTCTGGATTTGACATCTGGTTTTCTTGTCTTTCAAGGATGGCAGC  
CATTTTGAACGGTATTTGGAATACAATCGTTTTATTACCATTGAAGCTTATGACCCTAGTCGGCATTATGATTTGCTGGTTAC  
CAATAACCCGATTCTATAAGAAGGAACAGACACCAGTCTATTATTTAAAAAATGACTTGGATATGGAGGATTTGGTAGCGATT  
CGCCAGTTATTATCACTTAA

## 4106.7 (SEQ. ID. NO. 215)

ATGGAATTTTCAAAGAAAACACGTGAATTGTCAATTAATAAAAAATGCAGGAACGTACCCTGGACCTCTTGATTATCGGTGGAG  
GAATCACAGGAGCTGGTGTAGCCTTGCAGGCGGCAGCTAGCGGTCTTGAGACTGGTTTGATTGAAATGCAAGACTTTGCAGA  
AGGAACATCTAGTCGTTCAACAAAAATTTGGTTACGGAGGACTTCGTTACCTCAAAACAATTTGACGTAGAAGTGGTCTCAGAT  
ACGGTTTCTGAACGTGCAGTGGTTCAACAAATCGCTCCACACATTCCAAAATCAGATCCAATGCTCTTACCAGTTTACGATGA  
AGATGGAGCAACCTTTAGCCTCTTCCGCTTTAAAGTAGCCATGGACTTGTACGACCTCTTGGCAGGTGTTAGCAACACACCAG  
CTGCGAACAAGGTTTTGAGCAAGGATCAAGTCTTGGAAACGCCAACCAAACTTGAAGAAGGAAGGCTTGGTAGGAGTGGAG  
TGTATCTTGACTTCCGTAACAACGATGCGCGTCTCGTGATTGAAAACATCAAAACGTGCCAACCAAGACGGTGCCCTCATTGCC  
AACCACGTGAAGGCAGAAGGCTTCCCTCTTTCAGCAAAAGTGGCAAGATTACAGGTGTTGTAGCTCGTGATCTCTTGACAGACC  
AAGTGTTTGAAATCAAGGCCGCTCTGGTTATTAATACAACAGGTCTTGGAGTGATAAAGTACGTAATTTGTCTAATAAGGG  
AACGCAATCTCAAAATGCGCCCACTAAGGGAGTTCACTTGGTAGTAGATTCAAGCAAAATCAAGGTTTACAGCCAGTT  
TACTTCGACACAGGTTTGGGTGACGGTCGTATGGTCTTTGTTCTCCACGTGAAAACAAGACTTACTTTGGTACAACGTGATAC  
AGACTACACAGGTGATTTGGAGCATCCAAAAAGTAACCAAGAAGATGTAGATTATCTACTTGGCATTGTCAACAACCGCTTC  
CCAGAATCCAACATCACCATTGATGATATCGAAAGCAGCTGGGCAGGTCTTCG  
TCCATTGATTGCAGGGAACAGTGCCTCTGACTATAATGGTGAAAATAACGGTACCATCAGTGATGAAAGCTTTGACAACCTTG  
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CATCTGAGAAACATTTGGATCCATCTGCAGTTTCTCGTGGGTCTAGCTTGGACCGTGATGACAATGGTCTCTTGACTCTTGCT  
GGTGGTAAAAATCACAGACTACCGTAAGATGGCTGAAGGAGCTATGGAGCGCGTGGTTGACATCCTCAAGCAGAAATTTGAC  
CGTAGCTTTAAATGATCAATTTCTAAAACCTTACCCTGTTTCAGGTGGAGAATTGAACCCAGCAAAATGTGGATTGAGAAATCG  
AAGCCTTTGCGCAACTTGGAGTATCACGTGGTTTGGATAGCAAGGAAGCTCACTATCTGGCAAAATCTTACGGTTCAAATGCG  
ACCGAAAGTCTTTGCACTTGTCTCACAGCTTGAACAAGCGCCAGGACTCAGTTGGCAGATACTTTGTCCTTCACTATGCAA  
TGCGCAATGAGTTGACTCTTACGCCAGTTGACTTCTTCTCGTGTACCAATCACATGCTCTTTATGCGTGATAGCTTGGATA  
GTATCGTTGAGCCAATTTTGGATGAAATGGGACGATTCTATGACTGGACAGAAGAAGAAAAAGCAACTTACCGTGCTGATGT  
CGAAGCAGCTCTCGCTAACAACGATTTAGCAGAATTAATAAAATTA

## 4106.8 (SEQ. ID. NO. 216)

ATGATGAATGAATTAATTTGGAGAAATTTCTAGGGACTTTAATCCTGATTCTTCTAGGAAATGGTGTGTTGTCAGGTGTGGTTCT  
TCCTAAAACCAAGAGCAATAGCTCAGGTTGGATTGTGATTACTATGGGTTGGGGATTGCAGTTGCGGTTGCAGTCTTTGTAT  
CTGGCAAGCTCAGTCCAGCTTATTTAAACCCAGCTGTGACCATCGGTGTGGCCTTAAAGGTTGGTTTGCCTTGGGCTTCCGTT  
TTGCCTTATATCTTACGCCAGTTTCGACAGGGGCCATGCTGGGTGAGATTTTGGTTTGGTTGCAATTCAAACCTCACTATGAGGC  
AGAAGAAAAATGCAGGCAATATCCTGGCAACCTTCAGTACTGGACCAAGCATCAAGGATACTGTATCAAACTTGATTAGCGAA  
ATCCTTGGAACTTTTGTGTTGGTGTGACAATCTTTGCTTTGGGTCTTTACGACTTTCAGGCAGGTATCGGAACCTTTGCAATG  
GGAACCTTTGATTGTGCGGTATCGGTCTATCACTAGGTGGGACAACAGGTTATGCCTTGAACCCAGCTCGTGACCTTGGACCTCG  
TATCATGCACAGCATCTTGCCAATTCAAACAAGGGAGACGGAGACTGGTCTTACGCTTGGATTCTGTTGTAGGCCCTGTAT  
TCGGAGCAGCTTGGCAGTGCTTGTATTCTCACTTTTCTAG

## 4106.10 (SEQ. ID. NO. 217)

ATGAAAAGGACCTGGAGGAACCTATTCTGTGACAAATCTTAATACACCTTTTATGATTGGCAATATTGAGATTCCCAATCGTAC  
CGTTTTAGCGCCTATGGCTGGCGTGACCAACTCAGCCTTTCTGACTATCGCAAAAGGAGCTCGGAGCTGGACTCGTTGTAATGG  
AAATGGTCTCTGACAAGGGAATCCAATACAACAACGAAAAAACCTGCACATGCTTCATATCGATGAGGGCGAAAAACCTTG  
TCTCTATCCAACCTTTTGGTAGCGATGAAGACAGCCTAGCACGCGCAGCAGAAATTCATCCAAGAAAAACCAAGACCGGATAT  
CGTCGATATCAACATGGGCTGCCCTGTCAACAAAATCGTGAAGAACGAAGCTGGTGCTATGTGGCTCAAGGATCCAGACAAG  
ATTTACTCCATCACTCAACAGGTCCAGTCTGCTTGTATCCCACTACTGTCAAAAATGCGTACCGGCTGGGCGGACCCATC  
TCTTGACGTAGAAAATGCTCTCGCTGCTGAAGCTGCAAGTGTCTTGCCTCGCCATGCATGGCCGTACCCGTGAACAAATGT  
ATACTGGCCACGCAGACCTTGAGACCCTTTACAAGGTTGCCAAGCTCTAACCAAGATTCCATTATCGCCAACGGTGATATC  
CGTACTGTCCAAGAAAGCAACGCAATCGAAGAAGTTGGTGTGACGCAGTCATGATTGGCCGAGCTGCCATGGGAAAT  
CCTTACCTCTTCAACCAATCAACCATTACTTTGAAACAGGAGAAATCCTACCTGATTTGACCTTTGAAGACAAGATGAAGAT  
CGCCTACGAACACTTGAAACGATTGATTAACCTCAAAGGAGAAAAACGTGCGAGTTCGTGAATTCGCGGGTCTCGCTCCTCAC  
TATCTCCGTGGAACATCTGGCGCTGCCAACTCCGTGGAGCCATTTGCAAGCCAGCACCTTGGCAGAGATTGAAACCTCT  
TGCAATTGGAGAAGGCTTAA

## 4107.1 (SEQ. ID. NO. 218)

ATGACAAAGAAGAAAATTTGAGCGTATTTCTGTAATACACCGAGAAAAAGATTTTATGGCTCAAGTGGTATTTTCATGCGAGATA  
AAGAACAACCTAAGTATAGTGTCTTGAGCGTAAAATGTTTGATGCTGCTAAAAATCAAGATATGCTAGCTTATCAAAAAATA  
CGCAACTATCAAGCAGATAACAGATATTAGGGTACAAACAAGTGAGGCTGACATTTTAGAGGCTGTAAAAGAGGTTTATGTG  
TACAATCACATGAATGTTATCGGAGCTTGTGACGGGATATTATTTATCAGTCAATCACCAGCTTATGATAAGTTAAATAAGTG  
GTTTAATATCTATTCTGATTTGTATTTAGCGTTGTACCCTTGCCAAAAATGGGGTATATCATGAGATGGTAGGTATCTAG

## 4107.2 (SEQ. ID. NO. 219)

ATGAAAAATTTCAACGAGGCTGAGATGAAATTAATCTTATACTGATATTCGGACTTCTTTGACAGAAATTTCAACAAGAGAGG  
CAGAAGAGCTAGTTGCAGCTGGCAAGCGGGTCTTCTACATTGCCCCCACTCTTTCTTTTGAAGGAACGCGCGCTGCTG  
GAATACTTGTCCAGCAGGCTTCTTTTTCGATTACCGTCACGCGCTTGTCTCAAAATGGCTCGCTATCTGGTCTTGAATGATTTA  
CCAGCTAAAACTACTCTTGATGATATCGGTCTTGGGTGGCCTTTTACAAATGCCTTGGCGAACTCGATCCCAAGGACTTGGC

TGTTTATGGCGCTATTAAGCAGGATCCTCAATTGATCCAGCAGTTAATTGAGCTTTACCATGAGATGACCAAATCTCAGATGA  
 GTTTTTTGGACTTGGAGAATTTAACAGATGAGGATAAGAGGGCGGATTTACTCTTGATTTTTGAGAAAAGTAACAGCCTATCTT  
 AATCAAGGTACAGTTAGCCCAGGAAAGTCAGTTGTCCCATTTGATTGAGGCTATTGAGAATGACAAGGTAAGTAGTGATTTTA  
 ATCAAATCGCCTTGGCTATTGACGGCTTTACTCGTTTTCTGCTGAGGAAGAGCGGGTTGTGGACTTACTTCACGGCAAAGGT  
 GTTGAGATTGTTATCGGGGCTTATGCTAGTAAGAAAAGCCTATACCAGTCCTTTTAGCGAGGGCAATCTCTACCAAGCCAGCGT  
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 ACCTGAGCTACAAGCATTTCGTATTCTCTTGGGGGATGTAGCTTCTTACCAGTTATCTCTCAAAACCATTTTTGACCAGTATC  
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 GCTCT

CTATCTTGCCTCAAGCAGGATTTTACCTGTCTACGTCGGGTTTGACAGAGTTTTATCGCAATGAATACAGTTATTTCTACGCT  
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 CGACCAATCCAGCCATCGAAACCATCAAGAAAGAAAGCAAAATTTGGTGGAAAAAGACCAAGCCTTTATCAATTAGACAAT  
 GGACGCACTGCTTTGTGACGAGGCAAGGTGGACCGGATTGACGTTTGAAGCTAATGGAGCGATAGGATGAGTAC  
 AAATCCAGTCTGACTCAGTTCCAGTTTCTCATTTCTTAAATGGGCTCAATTCTCAGTTACCAACCTATCTTGTGCTCCCTAAAA  
 AGAGAAGGGGAGCAGAATTTTTCGGCGCCATGTACTTGGAAATGGCTGAACCTGTCCATCTCTGATGGCGGTAAAAAGTCT  
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 AATTTTATAACAAAAACAGGCTAATCAACTGACAGATGAGGAATTTTCAAGTCTTCTTGGACTACAATGCCTATCTTTACAA  
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 CAGCAACATCAGGCTATTACAGGCTTTGAAGCCAATTACCATCTGGGCAAGCCCGTTTCTAGAAAAGTTGGACCTAGCTG  
 ATGGCAAGCGTCTGGTCGGAGAAAACTCAAGCAAGCTTGGCTTGAAAAATAA  
 GAGAGGAGTTGAATCGATGA

4107.3 (SEQ. ID. NO. 220)

ATGAAGCTTATTCCTTTTAAAGTGAGGAGGAGATTCAAAAACTGCAAGAAGCAGAAGCAAAATTCGAGCAAGGAACAGAAG  
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 TTGTCATGGCAGAGCGCAATTCTGGACCAATTGGCGCGTGGTGTCGAAATTTCTCAACTCTTTATCTCAACCTTTACCGTCAAG  
 GCTGCAACTGAAATTAAGAACGTTTGAAGAAAAACGTAAGCAAAAAATCCAGAAACAGATGATGTCGACCTCAAAACA  
 CACTTGGGTGCGCAGTTGGCAGACCTACCAACGCTGCCATTGGAACCATGGATTCTTTCACACAAAAATTCCTTGGCAAAC  
 ATGGTTATCTGCTTGATATTGCACCTAATTTCCGTATTTTACAAAAACCAAGCGAGCAACTTATTCTCGAAAACGAAGCTTTT  
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 GTTAATCTGGATGAAATGAGAAAATCCCAACGCTATGCGCAAAAATCCCAAGAAAGGCAAGGCTCATTATCTCTCAAGAAAAAT  
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 CATGCTAGACACTCTTCTGTCTATTACAAATCCCTGCAAGACTACGCTTGGTTGCCCTTATGAAGTCTCCAATGTTTGGTTT  
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 TTCCTAGTCTGACCTATAGGCAGAACGAAGGAGTAATACAGCTACGAAGCTATTCTGAGCAGATGCGTTTGGTGTATGTTGCT  
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GGAAGTTTATAATACTCTTACCAGCGCAGCTATTGAACTTCCTAGTGTCAAACCCCAAGTCAAATCAAGAAATCTACGAAC  
CAGTTATGGATATGGAAGGTGTCGAGATTGCTGGTCAAGGTCAGTCAGTAGGCAAGAAATCAGCTTCGATTTGCCAGATTT  
TTCAACCAAGAAAGGTAAGTGGAGCTGAGATTGGTAGTGCTACTCACGAATCATGCGAGAGAATTGACCTCAGCCAGCA  
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TTCTGTTGACTACAAGACAGCCGCTATGATGAACCAAGTCAACTCGTAGACCCGCTATCGTGGTCAGTTAGCTCTATACGA  
AGAGGCTTTATCAGAGCCTATTGATTGAAAATATTGAAAAATACTTGATTTTACTCGGTAAAGACGAGGTTCAAGTTGTA  
AAAGTATAA

## 4109.1 (SEQ. ID. NO. 221)

ATGGAACCTGCTCGCCATGCTGAAAGCTTGGGAGTAGATGCTATTGCAACGATTCCACCAATTTATTTCCGCTTGCCAGAATA  
CTCAGTTGCCAAATACTGGAACGATATCAGTTCTGAGCTCCAAACACAGACTACGTGATTTACAACATTCCTCAATTTGGCAG  
GGGTGCTTTGACTCCAAGCCTTTACACAGAAATCTTGAAAAATCCTGTTATCGGTGTGAAGAACTCTTCTATGCCAGT  
CAAGATATCCAAACCTTTGTCAGCCTTGGTGGAGAAGACCATATCGTCTTAAATGGTCCTGATGAGCAGTTCCTAGGAGGAC  
GCCTCATGGGGCTAGGGCTGGTATCGGTGGTACTTATGGTGCTATGCCAGAAGCTCTTCTTGAAAATCAATCAGTTGATTGCG  
GATAAGGACCTAGAAACAGCGCGTGAATTGCAATGCTATCAACGCAATCATTGGTAACTCACTTCTGCTCATGGAAATA  
TGTACGGTGTATCAAGAAGTCTTGAAAATCAATGAAGGCTTGAATATTGGATCTGTTCTGTTACCATTTGACACCAGTGACT  
GAAGAAGATCGTCCAGTTGTAGAAGCGGCTGCTGCCTTGATTTCGTGAAACCAAGGAGCGCTTCTCTAA

## 4110.2 (SEQ. ID. NO. 222)

ATGTATAAGACAAAGTGTTCACGAGAGAAGTTAGTATTATTTTTAAAAATTTTCTTCCCAATCCTGATCTACCAATTTGCCAA  
TTATCTGCCTCTTTTGTGATACTGCAATGACAGGTCATAACAACACTATGGACTTGGCTGGTGTATCTATGGCAACCAGTA  
TCTGGAATCCTTTTACTACATTTCTAACAGGGATTGTGTCAGCCTTGGTGCCTATCATTGGTACCATTGTTGCGAGGCAAAA  
AGGAAGAAGTTGCGTCTGATTTTACCAATTTATTTATTTGGCCTTGGGCCTATCTGTGGTCTTGTGGGGATGGTACTTTCT  
TGGCACCAATAATCTTGAATCATATTGGGTTAGAAGCAGCAGTAGCGGCAGTAGCGGTTCGCTATCTTTGGTTTTATCTATC  
GGGATATCCCCCTTGTGCTCTTTAGCGTCATTGCTTCTTGGTGGATTGCTGGGCTTGACCAAACTGTCCATGTACCTCATG  
CTTTTGTACTCCCTCAAGATAGCGGATTAACTATCTTGATTATGGCTAAGTTTTCGCCCTTGATTATAGCTACCAAGTCA  
GCTGGTTTGAAGACATCCTTGGCCTACTGGGCTTGTGCTGGGATTCTGTTCTGGTTTTATTTAAACAGGAGAAGCTCAAAGC  
CTTACACCTTGAGAAACGAATTCACCTTAATATGGATAAAATTAAGGAAGGAGTTCGTTTAGGTCTGCCTATTGGGGAACT  
GTCTTCGCGGAAGTGGCTATCTTTTCAAGTCTTATGTACGCTTTCTATGAGTATCTCATCGCTATGGCTATTGTGCTTCTATGAAGTGGGA  
GCCAAGCGATTGATGATGCGAAAACCTATATTGGTCTAGGAAGATGGACTGCCCTCATTTTTGCGGCCCTTACCTTAACCTT  
CCTTTACATTTTTAGGGGAAATGTGGCCAGTCTTTATGGTAACGACCCAAAATTTATCGATTGACAGTGCGTTTTTTAACTTA  
TAGTCTTTTCTTCCAGTTAGCAGATACCTTTGCGGCGCGCTTCAGGGAATTTTGGCGGGGTATAAGGATACAGTTATTCCTTT  
TTACCTTGGTTTGGTTGTTATTTGGGGCGTAGCAATCCCTGTGTACGCTATTGA

## 4112.2 (SEQ. ID. NO. 223)

ATGAGTACTTTAGCAAAAATAGAAGCGCTCTTGTGTTAGAGCGGTGAAGATGGGATTCGGGTCCGCCAGTTAGCTGAAGTCC  
TCTCTGTCACCGACAGGCATCCAGCAAAAGTTTAGGAAAATTAGCCAGAAAGTATGAAAAGGACCCAGATTCCAGTTTGGC  
TTTGATTGAGACAAGTGGTGTATAGATTGGTGACCAAGCCTCAATTTGCGAGAGATTTGAAGGAATACTTAAGGCGCCT  
ATCAACCGAGCTTGTCTCGGGCTGCCCTTGAGACCTTGTCCATTATTGCCTACAAACAGCCGATTACGCGGATAGAAATTGA  
TGCCATCCGTGGAGTTAACTCGAGTGGAGCCTTGGCAAAAGTTGCAGGCTTTTGACCTGATAAAGGAAGACGGGAAAAAGGA  
AGTATTGGGGCGCCCCAACCTCTATGTGACTACGGATTATTTCCCTAGATTACATGGGGATAAACCATTTAGAAGAATTACCA  
GTGATTGATGAGCTTGAGATTCAAGCCCAAGAAAGCCAATTATTTGGTGAAAGGATAGAAGAAGATGAGAATCAATAA

## 4113.1 (SEQ. ID. NO. 224)

ATGGATACGATGATTAGTAGATTTTTTCGCCATTTATTTGAAGCCTTAAAAAGTTTGAAACGAAATGGTTGGATGACAGTAGC  
TGCTGTGATCTCAGTCAAGTATTACTTTGACCTTGGTGGCAATATTTGCATCTGTTATTTTCAATACAGCGAAACTAGCTACAG  
ATATTGAAAATAATGTCCGTGTAGTAGTTTATATCCGAAAGGATGTGGAAGATAATAGTCAGACAATTGAAAAAGAAAGGTCA  
AACTGTTACAAATAATGACTACCACAAGGTATATGATTCTTTGAAGAATGCTACCGTTAAAAGTGTACCTTTTCAAGTA  
AAGAAGAACAATATGAAAAATTAACCGAGATAATGGGAGATAACTGGAATAATCTTTGAAGGAGATGCCAATCCTCTCTATG  
ATGCCTATATTGTAGAGGCAAAACACTCCAAATGATGTAAAAACTATAGCCGAAGATGCTAAAAAAATGAAGGTGCTCTGA  
GGTTCAAGATGGCGGTGCCAATACAGAAAGACTCTTCAAGTTAGCTTCATTTATCCGTGTTTGGGGACTAGGGATTGCTGCTT  
TGTTAATTTTTATCGCAGTTTCTTGATTTCAAATACCATTGCTATTACCATTTTCCCGCAGTCGCGAAATTCAAATCATGC  
GCTTGGTCGGAAGCTAAAAACAGTTATATCCGTGGACCGTCTTGTGTTAGAAAGGAGCCTTTATCGGTTTATTGGGAGCTATCGCA  
CCATCTGTTTTGGTCTTTATTTGTTTATCAAATTTGTTTACCAATCTGTCAACAAATCGTTGGTAGGGCAAAATCTATCCATGATT  
AGTCCAGATTTATTTAGTCCGTTGATGATTGCCCTACTATTTGTGATTGGGGTTTTTCATTGGTTCATTGGGATCAGGAATATCC  
ATGCGCCGATTCTTGAAGATTAG

## 4117.1 (SEQ. ID. NO. 225)

ATGAAGAAAGTAAGATTTATTTTTTAGCTCTGCTATTTTCTTAGCTAGTCCAGAGGGTGCAATGGCTAGTGATGGTACTTG  
GCAAGGAAACAGTATCTGAAAGAAGATGGCAGTCAAGCAAGCAATGAGTGGGTTTTGATACTCATTATCAATCTTGGTTC  
TATATAAAAGCAGATGCTAACTATGCTGAAAAATGAATGGCTAAAGCAAGGTGACGACTATTTTACCTCAAATCTGGTGGCT  
ATATGGCCAAATCAGAATGGGTAGAAGACAAGGGAGCCTTTTATTATCTTGACCAAGATGGAAGATGAAAAGAAATGCTT  
GGGTAGGAACCTCCTATGTTGGTGCAACAGGTGCCAAAGTAATAGAAGACTGGGTCTATGATTCTCAATACGATGCTTGGTT  
TTATATCAAAAGCAGATGGACAGCAGCAGAGAAAGAAATGGCTGCAAAATTAAGGGAAGGACTATTATTTCAAATCCGGTGG  
TTATCTACTGACAAGTCAGTGGATTAATCAAGCTTATGTGAATGCTAGTGGTGCCAAAGTACAGCAAGGTTGGCTTTTGGACA  
AACAATACCAATCTTGGTTTTACATCAAGAAAAATGGAACATGCTGATAAAGAAATGGATTTTCGAGAAATGGTCACTATTA  
TTATCTAAAAATCCGGTGGCTACATGGCAGCAATGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATTTGATGGGA  
AAATGGCTGAAAAAGAAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTACATGACAGCCAA  
TGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATCTGATGGGAAAAATAGCTGAAAAAGAAATGGGTCTACGATTCT  
CATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTACATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTA

CCTCAAATCTGATGGGAAAAATAGCTGAAAAAGAAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCTGGT  
 GGCTACATGGCGAAAAATGAGACAGTAGATGGTTATCAGCTTGGGAAGCGATGGTAAATGGCTTGGAGGAAAAACTACAAAT  
 GAAAAATGCTGCTTACTATCAAGTAGTGCCCTGTTACAGCCAATGTTTATGATTTCAGATGGTGAAAAAGCTTTCCTATATATCGCA  
 AGGTAGTGTCTGATGGCTAGATAAGGATAGAAAAAGTGATGACAAGCGCTTGGCTATTACTATTTCTGGTTTGTGAGGCTAT  
 ATGAAAAACAGAAGATTACAAAGCGTAGATGCTAGTAAAGGACTTTATCCCTTATTATGAGAGTGATGGCCACCGTTTTTATCA  
 CTATGTGGCTCAGAATGCTAGTATCCAGTAGCTTCTCATCTTTCTGATATGGAAGTAGGCAAGAAATATTATTCGGCAGATG  
 GCCTGCATTTTGTATGGTTTAAAGCTTGAGAATCCCTTCTTTTCAAAGATTAAACAGAGGCTACAAACTACAGTGCTGAAGAA  
 TTGGATAAGGTATTTAGTTTGTCTAAACATAACAATAGCCTTTTGGAGAACAGGGCGCTACTTTTAAAGGAAGCCGAAGAAC  
 ATTACCATATCAATGCTCTTTATCTCCTTGCCCATAGTGCCCTAGAAAAGTAACCTGGGGAAGAAAGTAAAAATTGCCAAAGATAA  
 GAATAATTTCTTTGGCATTACAGCCTATGATACGACCCCTTACCTTTCTGCTAAGACATTTGATGATGTGGATAAGGGAATTT  
 TAGGTGCAACCAAGTGGATTAAAGGAAAAATTATATCGATAGGGGAAGAACTTTCTTGGAAACAAGGCTTCTGGTATGAATGT  
 GGAATATGCTTCAGACCCCTTATTGGGGCGAAAAAATTGCTAGTGTGATGATGAAAAATCAATGAGAAGCTAGGTGGCAAGA  
 TTAG

## 4119.2 (SEQ. ID. NO. 226)

ATGAAAAAAGTATTACAAAAATATTGGGCATGGGCTTTTGTGGTCATCCCCCTTGTGTACAAGCAATTTTCTTCTATGTGCC  
 GATGTTTCAAGGAGCCTTTTACAGTTTACCAACTGGACAGGATTGACTTATAACTACAAATTTGTTGGCTTAAACAACCTTTA  
 AGCTCCTCTTCATGGATCCAAAATTCATGAATGCGATTGGCTTTACCGCAATCATTGCGATTGCCATGGTGGTTGGTGAGATT  
 GCACTCGGGATCTTCATTGCGCGTGTCTTGAATTCATAAAATCAAAGGCCAAACCTTCTTCCGTGCTTGGTCTTCTTCCAGCT  
 GTTTTATCTGGTTTGACAGTGGCTTTGATCTTCAAGCAAGTCTTCAACTACGGTCTTCCAGCGATTGGAAATGCCCTTCATATT  
 GAATTTTCCAAACCAGTCTTTAGGGACTAAGTGGGGAGCAATCTTTGCGGCTGTCTTTGTCTTCTTTGGCAAGGGGTGGC  
 TATGCCCATCATCATCTTCTAGCTGGTTTGAATCTATTCCAAGTACAGATTACAGAGGCAGCAAGGATTGATGGTGCGACTA  
 GCAAGAAATTTTCTGGAAACATTGAATTGCCTTACTGTACCAAGTGTCTCTATGGTCTTTATCCTAGCCCTAAAAGGTGGG  
 CTGACTGCCCTTTGACCAAGTCTTTGCCATGACCGGTGGTGGTCCAAACAATGCCACAACCTCACTTGGGCTCTTGGTTTATAA  
 CTATGCCCTTTAAAAACAACCAATTTCGGTTATGCCAATGCCATTGCCGTAATCTTGTCTTCTTAATTGTAGTGATTTCGATCAT  
 CCAATTGAGAGTATCTAAGAAATTTGAAATTTAA

## 4119.3 (SEQ. ID. NO. 227)

ATGATGAAACAAGATGAAAAAGCCCTGATTGGCAAATACATTCTATTGATTCTAGGATCGGTTCTGATTTTAGTGCCCGC  
 TCCTTGTCTACCCTCTTTAGTTTCTTCAAACCCACTAAGGATATTGTAGATAATTTCTTTGGCTTTCCAACCAACTTCACATGGG  
 ACAACTTTAGCCGTCTCTTAGCTGATGGGATTGGAGGCTATTATTGGAAGTCTGTCTCATCACTGTCTTGTCTTTACTTGCAG  
 TAATGATCTTTATCCCTATGGCAGCCTACTCCATCGCTCGCAATATGAGTAAAAGAAAAGCCCTTACCATCATGTATACCCTC  
 TTAATCTCGGAATCTTCGTACCTTTCCAAGTCATCATGATTCCGATTACGGTTATGATGAGTAAACTCGGTTTGGCTAATACC  
 TTTGGTTTGTATCTTGCTCTACTTGACCTATGCGATTCCACAGACCCCTTTCTCTATGTTGGCTATACAAAATCTCGATTCCA  
 GAAAGTCTGGATGAAGCAGCAGAGATCGATGGGGCTAATCAATTTACAACCTATTTCCGCATCATCTTCCCAATGATGAAAC  
 CGATGTCATGGCACAACCATGATCATCAATGCCCTTTGGTTCTGGAATGACTTCATGTTGCCACTCCTTGTCTTGAACCGGGAT  
 TCCAAAATGTGGACTCTGCCCTTGTTCCTAATCAACTACGACAGGCAATTTTCAACGACTACGGACCAAGCTTGGCTCTTTA  
 CGTGGTCCGCATTATCAGTATCACCATTGTCTATCTTCTTCCAACGCCATATCATTTACGGAATGAGCAACGGGGCAGTGA  
 AGTAA

## 4119.4 (SEQ. ID. NO. 228)

ATGAAAAAGTATTCTTCAGAAAAATGGGGGAGCATCCGATGCTGCTTCTTTTCTTAGCTATAGTACTGTTATATCCATTCTTGCA  
 CAAAATGGATGGGTCTTGTGGCTTCAGTAGGAATGTTTCTATTACTATTTTCTTTTGGCACTATCAGTCGATTTTATCCCAT  
 AAATCTTTCGATGATTTTGCAGTTTGTCTTTGTTGTTAGTGTCTTGCAGTGTCTTTGCCAGTTTGAAGATTTCCAAATTTG  
 TGAAGAAATTTAACTATGCTTTTCTTTACCCCAATATGCAGGTGTGGCATCAGAACCAGGGCAGAAAGTGACCTTCTTTAATCCT  
 AATTATTTATGGAATTTTGTGTTTCTGTATTATGATTGCTTTCTATCTGTTTACAACGACCAAGTTGAATTTGGTTGAAAGTA  
 TTCTGTGTGATTGCAGGCTTTGTTAATCTCTTTGGTTTGAACCTTTACTCAAAATCGAACTGCCTTCTCTGCTATTATCGTGGG  
 GCAAAATATCTATCTTTACGACTATTAAAACTGGAAGGCCCTTTGGCTTAGTATTGGGGTCTTCCGCTTGGTTTGGTTTCTT  
 CTCCTTCTAGTGATTGGGAGTTCGAATGGGTACTTTAGACTCTTCTATGGAAGAACGCATTTCTATCTGGGATGCTGGGAT  
 GGCCTTGTTTAAGCAAAATCTTTTGGGGTGAAGGGCCATTGACCTATATGAACTCTTATCCTCGGATACATGCTCCTTATC  
 ATGAACATGCCACAGTCTTTATATTGATACGATTGAGTTACGGAATTTGTGGGACTATTTATTAGTTTGTCTTCTGTTG  
 CTCCTGTTGCTTGTATGATGGATATGAGTCAGGAGTCGGGGAAACGTCGGATTATCGGCCCTTTATCTATCTTTCTTACAGTG  
 GTTGCTGTGCACGGAATTTTGAATTTGACTTGGCTCTCTTCTGGATTTCAGTCAGGCTTATTTTCTTGTCTAGTTATGTGCAGTATTCAT  
 TGGAGCATCGAATGTTGGTATCGGACATGACGGATTAA

## 4120.1 (SEQ. ID. NO. 229)

ATGTCAAAGATGGATGTTTCAGAAAAATCATTGCACCGATGATGAAGTTTGTGAATATGCGTGGCATTATAGCTCTAAAAGATG  
 GGATGTTTAGCAATTTTGGCATTGACAGTAGTTGGTAGTTTGTCTTGATTATGGGACAATTGCCGTTTCAAGGATTAAATAAG  
 AGCATTGCTAGTGTTTTGGAGCTAATTGGACAGAGCCGTTTATGCAAGTATATTACAGGAACCTTTGCTATTATGGGTCTAAT  
 TTCTTGTTTTCAATTGCCTATTCTTATGCTAAGAAATAGCGGAGTAGAGGCTTTACCAGCTGGAGTTCTATCTGTATCTGCATT  
 CTTTATTTTGTAAAGATCATCTTATATCCCTAAACAAGGTGAGGCGATTGGGGACGCTATTAGTAAAGTTTGGTTTGGAGGCC  
 AAGGAATATCCGTGCTATCATTATAGGTTTGGTAGTAGGAAGTATTATACCTTCTTTATAAAGAGAAAAATTTGTTATTAAAG  
 ATGCCAGAACAAGTCCACAAGCTATTGCCAAACAGTTTGAAGCAATGATTCCAGCATTGTGAATTTTCTTATCTTCTATGAT  
 TGTATATATTTAGCGAAGTCAATTGACTAATGGCGGAACATTATAGAAATGATTTATCTGCTATTCAAGTTCCGTTGCAAG  
 GTTTAACTGGATCTTTGTATGGTCTATTGGAATTGCACTTTTATATCATTTTGTGGTGGTTTGGTGGTAACTCGG  
 TAGTAAATGGAGTAGTGACAGCTCTGCTTTTATCTAATCTTGATGCTAATAAAGCTATGTTAGCCTCTGCTAATCTATCATT  
 GAAAATGGTGCACATATTGTTACTCAACAATTTTATGATTCAATTTTAAATCTATCAGGTTCCAGGGATTACGTTTGGTCTTGTA  
 GTTGCCATGCTTTTTCAGCAAAAATCAAAACAATACCAAGCCCTAGGAAAAGTTGCAGCTTTTCCAGCAATATTTAACGTAA  
 ATGAGCCAGTTGATTTTGGATTTCGGATTGTCATGAATCCAGTTATGTTTGTACCTTTCAITCTTGTCTGTACTGCGAGCTGT  
 GATAGTATATGGAGCTATTGCAACAGGTTTTCATGCAGCCATTCTCAGGGGTAACATTGCCTTGGAGTACACCAGCTATTTTAT  
 CAGGATTTTGGTGGGTGGATGGCAAGGAGTTATTACTAGCTGGTGATATTAGCGATGTCTACATTGGTTTATTTTCCATTCT  
 TTAAGTACAGGATCGTTTACCTTACCAAAATGAAATCAAAACAATCTTAG

## 4121.2 (SEQ. ID. NO. 230)

ATGAAGAAAAAGGACTTAGTAGACCAACTAGTCTCAGAGATCGAGACGGGGAAAGTCAGGACACTGGGAATATACGGTCAT  
GGAGCTTCAGGTAATCAACCTTTGCACAGGAATTGTACCAAGCTTTAGATTCTACTACAGTAAATTTGCTAGAGACAGATC  
CTTATATCACCTCAGGACGCCATCTGGTAGTACCAAGGACGCGCCAATCAAAAGGTGACAGCCAGTCTGCCAGTGGCGCA  
TGAAGTGGAGAGTTTGCAGAGAGATATCCTTgCTTGACGGCGGGTATGGATGTCTTGA

## 4122.1 (SEQ. ID. NO. 231)

ATGAAGAAAAGATACCTAGTCTTGACAGCTTTGCTAGCCTTGAGTCTAGCAGCTTGTTCAAGAAAAACAAAAATGAAG  
ATGGAGAACTAAGACAGAACAGACAGCCAAAGCTGATGGAAACAGTCGGTAGTAAGTCTCAAGGAGCTGCCAGAGAAA  
GCAGAAGTGGTCAATAAAGGTGATTACTACAGCATTCAAGGGAAATACGATGAAATCATCGTAGCCAACAAACACTATCCA  
TTGTCTAAAGACTATAATCCAGGGGAAAAATCCAACAGCCAAGGCAGAGTTGGTCAAACATCAAAAGCGATGCAAGAGGCA  
GGTTTCCTATTAGTGATCATTACAGTGGTTTTAGAAGTTATGAACTCAGACCAAGCTCTATCAAGATTATGTCAACCAAGA  
TGGAAAGGCAGCAGCTGACCGTTACTCTGCCCGTCTGGCTATAGCGAACACCAGACAGGCTTGGCCTTTGATGTGATTGGG  
ACTGATGGTGATTTGGTGACAGAAAGAAAGCAGCCCAATGGCTCTTGGATCATGCAGCTGATTATGGCTTTGTTGTCCGTTA  
TCTCAAAGGCAAGGAAAAAGGAAACAGGCTATATGGCTGAAGAATGGCACCTGCGTTATGTAGGAAAAGAGCTAAAGAAAT  
TGCTGCAAGTGGTCTCAGTTTGGAAAGAATACTATGGCTTTGAAGGCGGAGACTACGTCGATTAA

## 4125.6 (SEQ. ID. NO. 232)

ATGCGTAAATCTTAATTATTTTGTGCTACCAAGTTTTTTGACCATTTCAAAAAGTCGTTAGCACAGAAAAAGAGTCGTCTA  
TACTTCGAAAAGAAATTTATACCTTTCAATCTGACTTTGGTATTTATTTAGAGAAAAATTAAGTTCTCCCATGGTTTATGG  
AGAGGTTCCGTGTTATGCGAATGAAGATTTAGTAGTGAATCTGGGAAATGACTCCCAAAACAAGTTTCAAATAACCGAG  
TGGCGCTTAAATAAACAAGGAATTCAGTATTTAAGCTATCAATCATCAATTTATAGCTGCGGACAAACGATTTTATATGA  
TCAATCAGAGGTAACCTCCAACAATAAAAAAGTATGGTTAGAATCTGACTTTAACTGTACAATAGTCCTTATGATTTAAAA  
GAAGTGAATCATCCTTATCAGCTTATTCGCAAGTATCAATCGACAAGACCATGTTGTAGAAGGAAGAGAATTTCTACATA  
TTGATCAGGCTGGATGGGTAGCTAAAGAATCAACTTCTGAAGAAGATAATCGGATGAGTAAAGTTCAAGAAATGTTATCTGA  
AAAAATATCAGAAAGATCTTTCTATTTATGTTAAGCAACTGACTACTGGAAAAGAGCTGGTATCAATCAAGATGAAAAG  
ATGTATGACGCCAGCGTTTGAAGCTCTTATCTCTATTATACGCAAGAAAAAATAAATGAGGGTCTTTATCAGTTAGATAC  
GACTGTAAATACGTATCTGCAGTCAATGATTTTCCAGGTTCTTATAAACCAGAGGGAAGTGGTAGTCTTCTAAAAAAGAA  
GATAATAAAGAATATTCTTTAAAGGATTTAATTACGAAAGTATCAAAAGAATCTGATAATGTAGTCTATAATCTATTGGGAT  
ATTACATTTCAAAACCAATCTGATGCCACATTCAAATCCAAGATGTCTGCCATTATGGGAGATGATTTGGGATCCAAAAGAAAA  
ATTGATTTCTTCTAAGATGGCCGGGAAGTTTATGGAAGCTATTTATAATCAAAATGGATTTGTGCTAGAGTCTTTGACTAAAA  
CAGATTTTGATAGTCAGCGAATTTGCCAAAGGTGTTTCTGTTAAAGTAG  
CTCATAAAATGGAGATGCGGATGAATTTAAGCATGATACGGGTGTTGTCTATGCAGATTCTCCATTTATCTTTCTATTTTCA  
CTAAGAATTCTGATTATGATACGATTTCTAAGATAGCCAAGGATGTTTATGAGGTTCTAAAAATGA

## 4125.7 (SEQ. ID. NO. 233)

ATGAAAAAACAAAATAATGGTTTAAATAAAAATCCTTTTCTATGGTTATTATTTATCTTTTCTTGTGACAGGATTCCAGTAT  
TTCTATTCTGGGAATAACTCAGGAGGAAGTCAGCAATCAACTATACTGAGTTGGTACAAGAAATACCGATGGTAATGTAA  
AAGAATTAACCTACCAACCAAAATGGTAGTGTTATCGAAGTTTCTGGTGTCTATAAAAAATCCTAAAAACAAGTAAAGAGAAAC  
AGGTATTCAGTTTTTACGCCATCTGTTACTAAGGTAGAGAAATTTACCAGCACTATTCTTCTGCAGATACTACCGTATCAG  
AATTGCAAAAACCTTGCTACTGACCATAAAGCAGAAGTAACTGTTAAGCATGAAAGTTCAAGTGGTATATGGATTAAATCTACT  
CGTATCCATTTGTGCCATTTGGAATTCTATTCTTCTCTATTCTATGATGGGAAATATGGGAGGAGGCAATGGCCGTAATC  
CAATGAGTTTTGGACGTAGTAAGGCTAAAGCAGCAAAATAAAGAAGATATTAAGTAAGATTTTCAGATGTTGCTGGAGCTGA  
GGAAGAAAAACAAGAACTAGTTGAAGTTTGAAGTTCTTAAAAAGTCCAAAACGATTTCACAAAACCTGGAGCCGATTTCGA  
GCAGGTGTTCTTTGGAGGGACCTCCGGGGACAGGTAACCTTTGCTTGCTAAGGCAGTCGCTGGAGAAGCAGGTGTTCCAT  
TCTTTAGTATCTCAGGTTCTGACTTTGTAGAAATGTTTGTGCGAGTTGGAGCTAGTCGTGTTCCGCTCTCTTTTGGAGGTGCCA  
AAAAAGCAGCACCAGCTATCATCTTTATCGATGAAATTTGATGCTGTTGGACGTCAACGTGGAGTCGGTCTCGGCGGAGGTAA  
TGACGAACGTGAACAAACCTTGAACCACTTTTGTAGTATGATGGTGGTTTGAAGGAAATGAAGGGATTATCGTCATCGCT  
GCGACAAACCGTTGAGATGTACTTGACCCTGCCCTTTTGCCTCCAGGACGTTTGTATAGAAAAGTATTGGTTGGTCTGCTCTGA  
TGTTAAAGGTGCTGAAGCAATCTTGAAAGTTACGCTATAAGAATAAGC  
CTTTAGCAGAAGATGTTGATTGAAATTAGTGGCTCAACAACTCCAGGCTTTGTTGGTGTGCTATTTAGAGAATGTCTTGAAT  
GAAGCAGCTTTAGTTGCTGCTCGTTCGCAATAAATCGATAATTGATGCTTCAGATATTGATGAAGCAGAAGATAGAGTTATTG  
CTGGACCTTCTAAGAAAGATAAGACAGTTTCACAAAAGAACGAGAATTGGTTGCTTACCATGAGGCAGGACATACCATTGT  
TGGTCTAGTCTTGTGCAATGCTCGCGTTGTCCATAAGGTTACAATTGTACCACGCGCGCTGCAGGCGGATACATGATTGCAC  
TTCTAAAGAGGATCAAAATGCTTCTATCTAAAGAAGATATGAAAGAGCAATTGGCTGGCTTAATGGGTGGACGTGTAGCTGA  
AGAAATTATCTTTAATGTCCAAACCACAGGAGCTTCAAACGACTTTGAACAAGCGACACAAATGGCACGTGCAATGGTTACA  
GAGTACGGTATGAGTGAAAAAATTTGGCCAGTACAATATGAAGGAAACCATGCTATGCTTGGTGCACAGAGTCTTCAAAAAAT  
CAATTTCAAGAACCAACAGCTTATGAAATTGATGAAGAGGTTCTGTTTATTAAATGAGGCACGAAATAAAGCTGCTGAAAT  
TATTCAGTCAAAATCGTGAAGCTCAAGTTAATTGCAGAAGCATTATTGAAATACGAAACATTGGATAGTACACAAATTA  
GCTCTTTACGAAACAGGAAAGATGCCTGAAGCAGTAGAAGAGGAATCTCATGCACTATCCTATGATGAAGTAAAGTCAAAA  
ATGAATGACGAAAAATAA

## 4125.10 (SEQ. ID. NO. 234)

ATGAGGGAACGAGATTTTTTAAATCATTTTCTCAAGAAGGGATATTTCAAAAAGCATGCTAAGGCGGTTCTAGCTCTTTCTGG  
TGGATTAGATTCATGTTTCTATTTAAGGTATTGTCTACTTATCAAAAAGAGTTAGAGATTGAATTCTAGCTCATGTGA  
ATCATAAGCAGAGAAATTGAATCAGATTGGGAAGAAAAAGGAATTAAGGAAGTTGGCTGCTGAAGCAGAGCTTCTATTTATAT  
CAGCAATTTTTCAGGAGAAATTTTCAGAAGCGCGTGCACGAAATTTTCGTTATGATTTTTTCAAGAGGTCATGAAAAAGACAG  
GTGCGACAGCTTTAGTCACTGCCACCATGCTGATGATCAGGTGGAACGATTTTATGCGCTTGATTGAGGAACCTCGTTG  
CGCTATGTCAGGAATTAAGGAGAAGCAAGTAGTCGGAGAGATGAAATTCATTCGTCCTTCTTGCATTTCAGAAAAAAG  
ACTTTCCATCAATTTTCACTTTGAAGATACATCAATCAGGAGAATCATTATTTTCGAAATCGTATTCGAAATTTCTACTTAC  
CAGAATTGGAAAAAGAAAATCCTCGATTAGGGATGCAATCTTAGGCATTGGCAATGAAATTTAGATTATGATTTGGCAAT  
AGCTGAATTATCTAACAATATTAATGTGGAAGATTACAGCAGTTATTTCTTACTCTGAGTCTACACAAAGAGTTTACTTC  
AACTTATCTGAATCGTTTTCAGATTGAAATCTTACAAAAGCTCAGTTTGTGTAAGTTTCAGCAGATTTTAAAAATCTAAAAGC  
CAGTATCGTATCCGATTAAAAATGGCTATGAATTGATAAAGAGTACCAACAGTTTCAGATTTGTAATAATCAGTCCGACAGG  
CTGATGAAAAGGAAGATGAACCTGTGTACACTATCAAAATCAGGTAGCTTATCAAGGATATTTATTTCTTTTGGACTTCCA

TTAGAAGGTGAATTAATTCAACAAATACCTGTTTCACGTGAAACATCCATACACATTTCGTCATCGAAAAACAGGAGATGTTT  
TGATTAATAAATGGGCATAGAAAAAACTCAGACGTTTATTATTG  
ATTTGAAAAATCCCTATGGAAAAAGAGAACTCTGCTCTTATTATTGAGCAATTTGGTGAAATTTGTCTCAATTTTGGGAATTGCG  
ACCAATAATTTGAGTAAAAAAACGAAAAATGATATAATGAACACTGTACTTTATATAGAAAAAATAGATAGGTAA

## 4126.1 (SEQ. ID. NO. 235)

ATGAAGCGTCTTCTCTTTAGTTAGAATGGTTATTTCCATCTTTCTGGTCTTTCTCATTCTCCTAGCTCTGGTTGGAACTTTCT  
ACTATCAATCAAGTTCTTCAGCCATTGAGGCCACCATTGAGGGCAACAGCCAAACGACCATCAGCCAGACTAGCCACTTTAT  
TCAGTCTTATATCAAAAACTAGAAACCACTCGACTGGTTTGACCCAGCAGACGGATGTTCTGGCCTATGCTGAGAATCCC  
AGTCAAGACAAGGTGCGAGGAATCCGAGATTTGTTTTGACCATCTTGAAGTCAGATAAGGACTTGAAAACCTGTTGTGCTGG  
TGACCAAACTCTGGTCAGGTCATTTCTACAGATGACAGTGTGCAGATGAAAACCTCCTCTGATATGATGGCTGAGGATTGGTA  
CCAAAAGGCCATTTCATCAGGGAGCTATGCCTGTTTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGGTCAATTTCTGCACTC  
AAGAAGTTGTTGATGCAAGGGAGCCAACTTGGTGCTCTCGTTGGATATTTCTTATGAAAACCTCTGGAAGCCTATCTCAAT  
CAACTCCAGTTGGGGCAGCAGGGCTTGCCTTCATTATCAATGAAAACCATGAATTTGTCTACCATCCTCAACACACAGTTTA  
TAGTTCGTCTAGCAAAATGGAGGCTATGAAACCCTACATCGATACAGGTCAGGGTTATACTCCTGGTCACAAATCCTACGTC  
AGTCAAGAGAAGATTGCAAGAACTGATTGGACGGTGTCTGGCGTGTGCATCATTGGAAAAAGTTAGACCAGGTTCCGGAGTCAGC  
TCTTGTGGACCTTGTCTGGGGCCAGTGTACATCTCTTCTGTCTGTCTGTCTAGTGTGGTTCAGTCTTAAACGCTGGATTG  
CTCCTTTGAAGGATTTGAGAGAAACCATGTTGGAAATTTGCTTCTGGTGTCTCAAAATCTTCGTGCCAAGGAAGTTGGTGCTAT  
GAAGTCAGAGAAGTAACTCGCAATTTAATGCTATGTTGGATCAGATTGATCAGTTGATGGTAGCTATTCGTAGCCAGGAAG  
AAACGACCCGTAAGAACTTCAAGCCCTTCGAGCCAGATTAATCCACATTTCTCTATAACACTTTGGACACCATCATC  
TGGATGGCTGAATTTTCATGATAGTCAGCGAGTGGTGCAGGTGACCAAGTCCTTGGCAACCTATTTCCGCTTGGCGCTCAATCA  
AGGCAAGGACTTGATTTGTCTCTCTGACGAAATCAATCATGTCCGCCAGTATCTCTTTATCCAGAAACAACGCTATGGAGATA  
AGCTGGAATACGAAATTAATGAAAATGTTGCCTTTGATAATTTAGTCTTACCCAAGCTGGTCTCAACCCCTTTGTAGAAAAAT  
GCTCTTTACCATGGCATTAAAGGAAAAAGGAAGGTGAGGGCCATATTAACCTTTCTGTCCAGAAACAGGATTCCGGGATTGGTCA  
TCCGATTGAGGATGATGGCGTTGGCTTCCAAGATGCTGGTGATAGTAGTCAAAGTCAAACGCTGGGGGAGTTGGTCT  
TCAAAATGTCGATCAACGGCTCAAACCTTCATTTTGAGGCAATTACCATATGAAGATTGATTCTAGACCCCCAAAAAGGGACG  
AAAGTTGAAATATATATAAATAGAATAGAACTAGCTAA

## 4126.7 (SEQ. ID. NO. 236)

ATGAAGCGTCTTCTCTTTAGTTAGAATGGTTATTTCCATCTTTCTGGTCTTTCTCATTCTCCTAGCTCTGGTTGGAACTTTCT  
ACTATCAATCAAGTTCTTCAGCCATTGAGGCCACCATTGAGGGCAACAGCCAAACGACCATCAGCCAGACTAGCCACTTTAT  
TCAGTCTTATATCAAAAACTAGAAACCACTCGACTGGTTTGACCCAGCAGACGGATGTTCTGGCCTATGCTGAGAATCCC  
AGTCAAGACAAGGTGCGAGGAATCCGAGATTTGTTTTGACCATCTTGAAGTCAGATAAGGACTTGAAAACCTGTTGTGCTGG  
TGACCAAACTCTGGTCAGGTCATTTCTACAGATGACAGTGTGCAGATGAAAACCTCCTCTGATATGATGGCTGAGGATTGGTA  
CCAAAAGGCCATTTCATCAGGGAGCTATGCCTGTTTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGGTCAATTTCTGCACTC  
AAGAAGTTGTTGATGCAAGGGAGCCAACTTGGTGCTCTGGTTGGATATTTCTTATGAAAACCTCTGGAAGCCTATCTCAAT  
CAACTCCAGTTGGGGCAGCAGGGCTTTGCCCTTCATTATCAATGAAAACCATGAATTTGTCTACCATCCTCAACACACAGTTTA  
TAGTTCGTCTAGCAAAATGGAGGCTATGAAACCCTACATCGATACAGGTCAGGGTTATACTCCTGGTCACAAATCCTACGTC  
AGTCAAGAGAAGATTGCAAGAACTGATTGGACGGTGTCTGGCGTGTGCATCATTGGAAAAAGTTAGACCAGGTTCCGGAGTCAGC  
TCTTGTGGACCTTGTCTGGGGCCAGTGTACATCTCTTCTGTCTGTCTGTCTAGTGTGGTTCAGTCTTAAACGCTGGATTG  
CTCCTTTGAAGGATTTGAGAGAAACCATGTTGGAAATTTGCTTCTGGTGTCTCAAAATCTTCGTGCCAAGGAAGTTGGTGCTAT  
GAAGTCAGAGAAGTAACTCGCAATTTAATGCTATGTTGGATCAGATTGATCAGTTGATGGTAGCTATTCGTAGCCAGGAAG  
AAACGACCCGTCAGTACCAACTTCAAGCCCTTCGAGCCAGATTAATCCACATTTCTCTATAACACTTTGGACACCATCATC  
TGGATGGCTGAATTTTCATGATAGTCAGCGAGTGGTGCAGGTGACCAAGTCCTTGGCAACCTATTTCCGCTTGGCGCTCAATCA  
AGGCAAGGACTTGATTTGTCTCTCTGACGAAATCAATCATGTCCGCCAGTATCTCTTTATCCAGAAACAACGCTATGGAGATA  
AGCTGGAATACGAAATTAATGAAAATGTTGCCTTTGATAATTTAGTCTTACCCAAGCTGGTCTCAACCCCTTTGTAGAAAAAT  
GCTCTTTACCATGGCATTAAAGGAAAAAGGAAGGTGAGGGCCATATTAACCTTTCTGTCCAGAAACAGGATTCCGGGATTGGTCA  
TCCGATTGAGGATGATGGCGTTGGCTTCCAAGATGCTGGTGATAGTAGTCAAAGTCAAACGCTGGGGGAGTTGGTCT  
TCAAAATGTCGATCAACGGCTCAAACCTTCATTTTGAGGCAATTACCATATGAAGATTGATTCTAGACCCCCAAAAAGGGACG  
AAAGTTGAAATATATATAAATAGAATAGAACTAGCTAA

## 4127.4 (SEQ. ID. NO. 237)

ATGTTTTTTAAATTAAGAGAAGCTCTTAAAGTCAAGCAGGTTTCGATCAAAAAATTTTATTTACAATTTTTATCGTTTTGGTC  
TTTCGATCGGAAGTACGATTACAGTTCCCTGGTGTGAATGCCAATAGCTTGAATGCTTTAAGTGGATTATCCTTCTTAAACAT  
GTTGAGCTTGGTGTGCGGGGAATGCCCTAAAAAACTTTTCGATTTTTGCCCTAGGAGTTAGTCCCTATATCACCGCTTCTATTGT  
TGTTCAACTCTTGCAATGGATATTTTACCCAAGTTTGTAGAGTGGGGTAAACAAGGGGAAGTAGGTGCAAGAAAAATTTGAAT  
CAAGCTACTCGTTATATTGCTCTAGTCTCGCTTTTGTGCAATCTATCGGGATTACAGCTGGTTTTAATACCTTGGCTGGAGCT  
CAATTGATTAAAACTGCTTTAACTCCACAAGTTTTTCTGACGATTGGTATCATCTTAACAGCTGGTAGTATGATTGTCACTTGG  
TTGGGTGAGCAAAATACAGATAAGGGATACGGAACGGTGTTCATGATTATCTTTGCCGGGATTGTTTCTCAATTCCAGA  
GATGATTACAGGCATCTATGTGGACTACTTTGTGAACGTCCCAAGTAGCCGTATCATTCTCATCTATTTCGTAATCATTTT  
GATTATTACTGTATTGTTGATTATTTACTTTACAACCTTATGTTCAACAAGCAGAATACAAAAATCCAAATCCAATATACTAAGG  
TTGCACAAGGTGCTCCATCTAGCTCTTACCTTCCGTTAAAAGTAAACCCTGCTGGAGTTATCCCTGTTATCTTTGCCAGTTTCA  
TTACTGCAGCGCTGCGGCTATTCTTCAGTTTTTGAAGTCCACAGGTGATGATTGGGCTTGGGTAAAGGTAGCACAAGAGATG  
TTGGCAACTACTTCTCCAAGCTGATTGCCATGATGCTTTGTGATTATCTCTTTACATTCTTCTATACGTTGTACAGATTA  
ATCCTGAAAAAGCAGCAGACGACCTACAAAAGAGTGGTGCTATATCCATGGAGTTCTGCTGGTAAAGGTACAGAGAATA  
TATGTCTAAACTTCTTCGTCGTCTTGAACCTGTTGGTTCCCTCTTCTTGGTGTGA

## 4127.5 (SEQ. ID. NO. 238)

ATGGATATTAGACAAGTTACTGAAACCATCGCCATGATTGAGGAGCAAACTTCGATATTAGAACCATTACCATGGGGATT  
CTCTTTTGGACTGTATCGATCCAGATATCAATCGTGTGCGGAGAAAACTCTACAAAAAATTACGACAAAGCGCGCTAATTT  
AGTAACTTGTGGTGATGAAATTCGCGCTGAGTTGGGAATTCCTATCGTTAATAAGCGTGTATCGGTGACACCTATTTCTCTGA  
TTGGGGCAGCGACAGATGCGACGGACTACGTGGTTCTGGCAAAAGCGCTTGATAAGGCTGCGAAAGAGATTGGTGTGGACT  
TTATTGGTGGTTTTTCTGCCTTAGTACAAAAAGGTTATCAAAAGGGAGATGAGATTCTCATCAATTCCATTCTCGCGCTTTG



GCTGAGACGGATAAGGTCTGCTCGTCAGTCAATATCGGCTCAACCAAGTCTGGTATTAATATGACGGCTGTGGCAGATATGG  
 GACGAATTATCAAGGAAACAGCAAACTCTTCAGATATGGGAGTGGCCAGTTGGTTGTATTGCTAATGCTGTTGAGGACAA  
 TCCATTTATGGCGGGTGCCTTTTCATGGTGTGGGGAAGCAGATGTTATCATCAATGTCGGAGTTTCTGGTCTGCTGTTGTGA  
 AACGTGCTTTGGAAAAAGTTCTGGGACAGAGCTTTGATGTAGTACCCGAAACAGTTAAGAAAACTGCCTTTAAAAATCACTCG  
 TATCGGTCAATTGGTTGGTCAAAATGGCCAGTGAGAGACTGGGTGTGGAGTTTGGTATTGTGGACTTGAGTTTGGCACCACCC  
 CCTGCGGTTGGAGACTCTGTGGCAGTGTCTTGGAGAAATGGGGCTAGAAACAGTTGGCAGCATGGAACGACGGCTGCCT  
 TGGCCCTCTTGAACGACCAAGTTAAAAAGGGTGGAGTGTGGCCTGCAACCAAGTCTGGTGGTTTATCTGGTGCCTTTATCCCT  
 GTTCTGAGGATGAAGGAATGATTGCTGCAGTGCAAAATGGCTCTCTTAAGTTTAAAGAAAACTAGAAGGATATGACGGCTAAT  
 GTTCTGTTGGATTGGATATGATTGCCATCCAGAGATACGCTGTGAAACTATTGCGGCTATGATTGCGGATGAAGCAGC  
 AATCGGTGTTATCAACATGAAACAAACAGCTGTTCGTATCATTCCTCAAGGAAAAAGGCGATATGATTGAGTTTGGTGGT  
 CTATTAGAACTGCACCCGTTATGAAGGTTAATGGGGCTTCGTCTGTCGACTTCTCTCGCGGTGGACAAATCCAGCACC  
 AATTCATAGTTTAAAAAATTA

## 4128.1 (SEQ. ID. NO. 239)

ATGACACAGATTATTGATGGGAAAGCTTTAGCGGCCAAATTGCAGGGCAGTTGGCTGAAAAGACTGCAAAATTAAGGAA  
 GAAACAGGTCTAGTGCTGGTTTGGTAGTGATTTTGGTTGGGACAATCCAGCCAGCCAAGTCTACGTTGCAACAAGGAGA  
 GGTGAGCCCTTGGCGGTGGTTTCCGTAGCGAAGTAGTACGGGTTCAGAGACCAATTAAGAGGAATTGTAGACCTGAT  
 TGCTAAATACAATCAGGATCCAGCTTGGCATGGGATTTTGGTTCAGTTGCCATTACCAAAACACATTGATGAAGAGGCGGTT  
 CTATTGGCTATTGACCCAGAAAAGGATGTGGATGGTTTCCATCCTCTAAACATGGGGCGTCTTGGTCTGGTCACTCCAGTCAT  
 GATTCCTTCGACACCGGCAGGAATTATGGAATGTTCCATGAATATGGGATTGACTTGAAGGTAATAATGCAGTCGTATC  
 GGTGATCCAAATATTGTGCGAAAACCTATGGCCAGCTTCTTTTGGCAAGAAATGCAACAGTAACCTTGACTCACTACGCTAC  
 TCATAACTCTTCAAGGTGGCTGCAAAAGCAGATATCTGGTTGTGCAATCGGTGCTGCCAAGTTTGTGACTGCTGACTTTG  
 TCAAACAGGTGGCGGTGACTTGGAGTTGGGATGAACCGCGATGAAAATGGTAAGCTCTGTGGGGATGTTGATTATGAGGC  
 GGTGCCCCACTTGCTAGCCACATTACGCCAGTCCCTGGAGGTGTGCGTCTATGACCATTACTATGCTGATGGAGCAACCT  
 ATCAGGCAGCACTTAGGACATTGGATAGAAAAATA

## 4128.2 (SEQ. ID. NO. 240)

ATGTCTAAATTTAATCGTATTCAATTTGGTGGTACTGGATTCTGTAGGAATCGGTGCAGCACCAGATGCTAATAACTTTGTCAA  
 TGCAGGGGTTCAGATGGAGCTTCTGACACACTGGGACACATTTCAAAAACAGTTGGTTTGAATGTCCCAACATGGCTAAA  
 ATAGGCTTTGGAAATATCTCTCGTGAAACTCCTTTAAGACTTGGCAGCTGAAAGCAATCCAACTGGGATATGACACAAAT  
 TAGAGGAAGTATCTCTTGGTAAGGATACTATGACTGGACACTGGGAAATCATGGGACTCAACATTACTGAGCCTTTGATAC  
 TTTCTGGAACGGATTCCAGAAAGAAATCCTGACAAAAATCGAAGAAATCTCAGGACGCAAGGTTATTCTGTAAGCCAACAAA  
 CCTTATTCAGGAACGGCTGTTATCTATGATTTTGGACCACGTCAGATGGAAACTGGAGAGTTGATTATCTATATACTTCACTG  
 CCCTGTTTTGCAGATTGCTGCCCACGAAGACATTATTCCTTTGGATGAATTGTACCGTATCTGTGAATACGCTCGTTTCGATTAC  
 CCTTGAGCGTCTGCCCCTTCTTGGTGCATCATTGCTCGCCCTTATGTAGGTGAACCAGGTAACCTCACTCGTACGGCAAAACC  
 GTCGTGACTTGGCTGTATCTCCATTTTCCCAACTGTTTTGGATAAAATGAATGAGGCTGGTATCGTACTTATGCTGTGGGT  
 AAATCAACGATATCTTTAACGGTGTCTGGTATCAACCAATGACATGGGTCAACAACAGTCAAATAGTCATGGAATTGATACACT  
 ATTGAAGACTATGGGACTTGTGAGTTTGAAGGATTCTCATTCACAAACCTAGTTGACTTTGATGCCCTTTACGGCCATC  
 GTCGTAATGCTACGGTTACCGTATGCTTGCATGAGTTGATGAACGCTTACCTGAAATATTCGCAGCTATGAGAGAGAAT  
 GACCTTCTTTGATTGCTGCGGACCATGGAATGACCAACGATATGCAGGAACGGATCACACTCGGGAATATATTCATTTGTT  
 GGCCTATAGCCCTGCCTTTAAAGGAAATGGTCTCATTCCAGTAGGACATTTTGCAGATATTTACGCGACTGTTGCCGATAACT  
 TTGGTGTGGAACCTGCTATGATTGGGGAAAGTTTCTTAGATAAAATGGTATAA

## 4129.2 (SEQ. ID. NO. 241)

ATGTTTATTTCCATCAGTGCTGGAATTGTGACATTTTACTAACTTTAGTAGAAAATCCGGCCCTTTATCCAATTTTATAGAAAG  
 GCGCAAATACAGGCCAGCAGATGCTAGGATGTCAAAACAGCATAGGCCAAAAGCTGGGACTCCTACAAATGGGAGGTTTG  
 GTTTCTTGATTACTTCTTGGTTGCTTCTTTTCCGCTATTAGTAGCCAATTCAGCAATAATGTGGGAATGATTTGT  
 TCATCTTGGTCTTGTATGGCTTGGTCCGATTTTTAGATGACTTTCTCAAGGTCTTTTGTAAAAATCAATGAGGGGCTTAATCCTA  
 AGCAAAAATAGCTCTTCAGCTTCTAGGTGGAGTTATCTTCTATCTTTCTATGAGCGCGGTGGCGATATCCTGTCTGCTTTG  
 GTTATCCAGTTCATTTGGGATTTTTCTATATTTCTTCTGCTCTTTCTGGCTAGTGGGTTTTTCAACGCAGTAACTTGACAGA  
 CGGTGTTGACGGTTAGCTAGTATTTCCGTTGTGATTGTTGCTGAGTATGGAGTTATTGCCATATGTGCAAGTATGATACACT  
 ATATTCTCTAGTGATTCTTGCCATGATTGGTGGTTTGTCTGGTTTCTTCTATCTTTAACCATAAGCCTGCCAAGGTCTTTATGG  
 GTGATGTGGGAAGTTTGGCCCTAGGTGGGATGCTGGCAGCTATCTCTATGGCTCTCCACCAAGAATGGACTCTCTTGATTATC  
 GGAATTGTGTATGTTTTTGAACCAACTTCTGTTATGATGCAAGTCAAGTTATTTCAAACTGACAGGTGGTAAACGTTATTTCCG  
 TATGACGCTGTACATCACCATTGAGCTTTGGGGATGTCTGGTAAAGGAAATCCTTGGAGCGAGTGGAAGGTTGACTTCT  
 TCTTTTGGGGAGTGGGACTTCTAGCAAGTCTCTGACCCTAGCAATTTTATATTTGATGTAA

## 4133.1 (SEQ. ID. NO. 242)

TTGTTTAAAGAAAAATAAAGACATTCTTAATATTGCATTGCCAGCTATGGGTGAAAACCTTTTTCAGATGCTAATGGGAATGGT  
 GGACAGTTATTGGTTGCTCATTTAGGATTGATAGCTATTTACGGGGTTTCAGTAGCTGGTAATATTATCACCATTTATCAGG  
 CGATTTTCATCGCTCTGGGAGCTGCTATTTCCAGTGTTATTTCAAAAAGCATAGGGCAGAAAGACCAGTCGAAGTTGGCCTAT  
 CATGTGACTGAGGCGTTGAAGATTACCTTACTATTAAGTTTCCCTTTAGGATTTTGTCCATCTTCGCTGGGAAAGAGATGAT  
 AGGACTTTTGGGGACGGAGAGGGATGTAGCTGAGAGTGGTGGACTGTATCTATCTTTGGTAGGCGGATCGATTGTTCTCTTA  
 GGTTTAATGACTAGTCTAGGAGCCTTGATTCGTGCAACGCATAATCCACGCTCTGCCTCTCTATGTTAGTTTTTATCCAATGCC  
 TTGAATATCTTTTCAAGTCTAGCTATTTTGTCTGGATATGGGATAGCTGGTGTGCTTGGGGACAATTTGTGCTCTCGT  
 TTGGTTGGTCTTGTGATTTTGTGGTCACAATTAACCTGCCTTATGGGAAGCCAACCTTTTGGTTTATAGATAAGGAAGTGTGAC  
 CTTGGCTTTACCAGCAGCTGGAGAGCGACTTATGATGAGGGCTGGAGATGTAGTATCATTGCCCTTGGTCTGTTCTTTTGGGA  
 CGGAGGCGAGTTGCTGGGAATGCAATCGGAGAAAGTCTTGACCCAGTTTAACTATATGCTGCTTTGGCGTCTGCTACGGCAAC  
 GGTGATGCTGTTGGCCGAGCAGTTGGAGAGGATGTTGAGAAAAGAGTTGCTAGTTTGAAGTAAACAAACCTTTTGGCTTCT  
 CTGTTCTCATGTTGCCCTGTCTTATGATATATGCTTGGGTGTACCATTAACCTCATCTCTATACGACTGATTCTCTAGCG  
 GTGGAGGCTAGTGTCTAGTGACACTGTTTCACTACTTGGGACCCCTATGACGACAGGAACAGTCATCTATACGGCAGTCTG  
 GCAGGGAATGGAATGACAGCCTCCCTTTTATGCGACAAGTATAGGAATGTGGTGTATCCGCAATGGGACAGGATATCTG  
 ATGGGGATTGTGCTTGGTTGGGGCTTGCCTGGTATTTGGGCAGGGTCTCTCTTGGATAATGGTTTTGCTGGTTATTTCTACGC  
 TATCGTTACCAGCGCTATATGAGCTTGAAAGGATAG

## 4135.2 (SEQ. ID. NO. 243)

ATGCAAACCTCAAGAAAAACACTCGCAAGCAGCCGTTCTTGGCTTGCAGCACTTACTAGCCATGTACTCAGGATCTATCCTGG  
 TTCCCATCATGATTGCGACAGCCCTTGGCTATTGAGCTGAGCAGTTGACCTACCTGATTCTACAGATATCTTCATGTGTGGG  
 GTGGCAACCTTCTCCTCAACTCCAACCTCAACAACTACTTTGGGATTGGACTCCCAGTCTTCTTGGAGTTGCATTCCAGTCGGT  
 CGCTCCCTTGATTATGTTGGGCAAGCCATGGTAGTGGCGCTATGTTTGGTGGCCCTTATCGCATCTGGGATTTACGTGGTTC  
 TTGTTTCAGGCATCTTCTCAAAAGTAGCCAATCTCTTCCCATCTATCGTAACAGGATCTGTTATTACCACGATTGGTTAACT  
 TGATCCCTGTGCGTATTGGAAATATGGGAAATAACGTTCCAGAGCCAACCTGGTCAAAAGTCTCTTGGCTTGCAGCTATTACTGTT  
 CTGATTATCCTCTTGATCAACATCTTTACCAAAGGATTTATCAAGTCTATCTCTATTTTGGATTGGTCTGGTGTGGAACTGCC  
 ATTGCTGCTACTATGGGCTTGGTGGACTTCTCTCTGTTGCGGTAGCTCCACTTGTCCATGTCCCAACTCCACTCTACTTTGGG  
 ATGCCAACCTTTGAAATCTCATCTATTGTCTATGATGTGTATCATCGCAACGGTGTCTATGGTTGAGTCAACTGGTGTATCTG  
 GCCTTGTCTGATATCACAAGGATCCAATCGACAGCAGCGCCCTTCGCAACGGATACCGCGCAGAAGGTTTGGCCGTACTTC  
 TCGGAGGAATCTTTAACACCTTCCCTTACACCGGATTTTCAAAAAACGTTGGTTTGGTTAAATTTGTCAGGCATCAAAAAACGC  
 CTGCCAATCTACTACGCAGCTGGTTTCTTGGTCTCTCTTGGACTGCTTCTAAGTTTGGCGCCCTTGGCCAAATCATTCCAAGC  
 TCCGTCTCGGTGGTGCCATGCTGGTAATGTTTGGTTTGTATCAATTCAAGGGATGCAAAATCCTCGCCCGTGTGACTTTGCT  
 AACATGAACACAACCTCTTATCGCAGCTGTTTCAATCGCTGCAGGTGTGGTCTCAACAACAGTAATCTCTTGTGACAT  
 GCCGACAGCCTTCCAATGTTCTTCTCAACCGGAATCGCTAGCCAGCCTACTCGCTATTGTCTCAATGCCGATTAAATC  
 AAAAAAGAAATAA

## 4136.2 (SEQ. ID. NO. 244)

ATGAAAGATAGATAAAAAAGAAATTTACAAGACAAGGGAAAGGTGACTGTTAATGATTGGCTCAGGCTTTGGGAAAAGAC  
 AGTTCCAAGGATTTTCGTGAGTTGATTAAAACCTTGTCTTAATGGAAAGAAAGCACCAAAATTCGTTTGAAGAAGATGGTA  
 GTCTGACATTAGAAATTAAGAAAAACATGAGATTACCTCAAGGGGATTTTCATGCCCATAAAAATGGCTTTGGCTTTGTT  
 AGTCTGGAAAGGCGAGGAGGACGACCTTTTGTAGGGAATAATGATGCAACTATGCTATTGATGGTGATACCGTCGAGGTAG  
 TGATTAAGAAAGTCGCTGACCGCAATAAGGGAAACAGCAGCAGAAGCCAAAATTATTGATATCCTAGAACACAGTTTGACAA  
 CAGTTGTGCGGCAAAATCGTTCTGGATCAGGAAAAACCTAAGTATGCTGGCTATATTCGTTCAAAAAATCAGAAAAACAGTCA  
 ACCGATTTATGTTAAGAAACAGCCCTAAAAATTAGAAGGAACAGAAGTTCTCAAAGTCTTTATCGATAAAATCCCAAGCAAG  
 AAACATGATTTCTTTGTCGCGAGTGTTCTCGATGTAGTGGGACACTCAACGGATGTGCGAATTGATGTTCTTGAGGTCTTGGA  
 ATCAATGGACATTGTATCCGAGTTTCCAGAAGCTGTTGTTAAGGAAGCAGAAAGTGTGCTGATGCTCCGTCTCAAAAGGAT  
 ATGGAAGGTGCTGATGATTAAGAGATGAAATACCTTTACCATTGACGGTGGCGATGCCAAGGACTTGGACGATGCAAGTGC  
 ATATACAGGCTCGAAAAATGGCAATCTGGAGTTTGGGTTTACATCGCAGATGTTTCTTATTATGTGACCGAGGGGTCTGCC  
 CTTGACAAGGAAGCCCTTAACCGTGGCAGTTCTGTTTACGTGACAGACCGAGTGGTGCCAATGCTTCCAGAACGACTATCAA  
 ATGGCATCTGCTCTCTCAATCCCAAGTTGACCGCTGACCCAGTCTGCTATTAT  
 GGAGATTGATAAACATGGTGTGTTGGTCAACTATACCATTACACAAACAGTTATCAAGACCAGTTTTCTGATGACCTATAGC  
 GATGTCAATGATATCCTAGCTGGCGATGAAGAAAAAGAGAAAAAATATCATAAAATTTGATCAAGTATCGAACTCATGGCC  
 AAGCTTCATGAACTTTAGAAAACATGCGTGTGAAACGTGGAGCTCTCAATTTTGATACCAATGAAGCGAAGATTTAGTGG  
 ATAAACAAGGTAAGCCTGTTGATATCGTTCTTCGGCAGCGTGGTATTGCCGAGCGGATGATTGAGTCTTTATGTTGATGGCT  
 AATGAAACAGTTGCCGAACATTTACGCAAGTTGGATTTGCTTTTATCTATCGAATTCACGAGGAGCCTAAGGCTGAAAAGG  
 TTCAGAAGTTTATTGATTATGCTTCGAGTTTGGCTTGGCATTATGGAAGTCCAGTGAGATTAGTCAGGAGGCACTTCAA  
 GACATCATGCGTGCTGTTGAGGGAGAACCTTATGCAGATGTTATGTCATGATGCTTCTCGCTCTATGCAGAGGCTCGTTA  
 TTCGGAGCACATACGGCCACTATGGACTAGCTGACTATTATCTACTTACCAGTCCAATTCGCTGTTTCCAGACCTTCCAGC  
 TTCTGTTTACCGTATGATTTCGGGATTACGGCCGTTCTAAGGAAATACGACAGCATTTTGAACAAGTGATTCCAGAGATTGCG  
 ACCAGTCTTCCAACCGTGAACGTGCTGCCATAGAAGCTGACCGTGAAGTCAAGCCATGAAAAAGGCTGAGTATATGGAA  
 GAATACGTGGGTGAAGAGATGATGTCAGTTGTATCAAGTATTTGCAAAATTCGGTCTCTTTGTCGAATTGCCAAACACAGTTGA  
 AGGCTTGATTACACTACTAATCTGCCTGAATTTATCAATTTACATGAGCGTGAATTTGACTCTTCGTGGAGAAAAATCAGGTA  
 TCACTTTCCGAGTGGGTGAGCAGATCCGTATCCGTGTTGAAAGAGCGGATAAAATGACTGGAGAGATTGATTTTTCATTCTGA  
 CCTAGTGAGTTTGATGTGATTGAAAAAGGCTTGAACAGTCTAGTCGT  
 AGTGGCAGAGGGCGTGATTCAAATCGTCTCGGATAAGAAGGAAGCAAGAGAAAAATCAGGACGCTCAAATGATAAGCGT  
 AAGCATTCACAAAAAGCAAGAAAGAAAAAGAAAGAAACCTTTTACAAGGAAGTAGCTAAGAAAGGAGCCAAGCATGG  
 CAAAGGGCGAGGGAAAGGTGCTCGCACAAAATAA

## 4137.2 (SEQ. ID. NO. 245)

ATGGGCACAACAGGATTTACAATAATTGACTTAATTATCTTGATTGTTTATTTACTTGGCGTGTGGTTGCAGGTATCTATTTC  
 TCTAAAAAAGAGATGAAAGGAAAAAGAGTTCTTTAAAGGAGATGGTTCGGTTCTTGGTATGTTACTTCGGTATCCATTTTTC  
 CACAATGCTCAGTCCGATTTCTTCTTGGGACTCGCTGGTAGCTCTTATGCAGGTAGCTGGATTTTATGGTGTGCTAATTAGG  
 GATGGTAGTAGCTATTCCACTGACAATTCGTTTATCTTACCTATCTTTGACGGATAGACATCGATACGGCATATGATTACTT  
 GGATAAACGTTTAAATTCTAAAGCACTTCGTATTATTTACGCACTTGTGTTATTTATCAATTGGGACGTATGCTATCAT  
 TATGTACCTCCCATCAGCTGGTTTATCAGTATTGACAGGAATTGACATCAATATTTTGAATTATTTGATGGGTGATGTTGCAAT  
 TGTTTATCTTATACTGGTGGTCTAAAAATCCGTATTATGGACAGCACTTTATTCAAGGTGTGATTCTGATTAGTGGTGTCTGTTT  
 AGCTTTATTTGTACTGATTGCTAATATTAAAGGTGGCTTTGGTGCAGTAGCAGAAACATTAGCAAACGGGAAATTCCTTGCTG  
 CAAATGAAAAACCTTTTCGATCCTAAGCTTGTCTTCAAACTCCATCTTTTAAATTGTGATGGGTTCAGGCTTTACAATCTTGTCTT  
 CCTATGCTTCACTCAAGATTGGTTCAACGTTTACTACAACACAAAAATTAAGAACTTAATAAGATGTTGTTCCAAAAAC  
 GGTGTTTGTGCACTTGCAACTGCAACAGTCTTTACTTGATTGGTACAGGCTTGTACGTATTCTCAAGTACAAAAATGCAGA  
 TAGTGCAGCTAGCAATATCCCTCAAGACCAAACTTTATGTACTTTATGCATACCAGTTACCAGTAGGTATCACAGGTTTGA  
 TCTTGGCAGCGGATTTATGCAGCATCTCAATCAACTATTTCAACAGGTTTGAACCTGTTGCAACTTTCATGGACATTTGGATATT  
 CAAGATGTCAATTTCTAAAAATATGTCAGACAATCGTCTGACGAAATTTGCAAAATTCGATATCTCTAGCAAGGTTTATCTC  
 AATTGGTGTGTTCCATTGTCTAGGCTCACTCAGATATTAATCTGCATACGAATGGTTCAATAGTTTTCATGGGACTTGTACTTG  
 GTCTACTTGGTGGTGTATTTATCTTGGATTGTTTCTAAAAAAGCAAAATAAACAAGGTGCTTATGCAGCGCTGATTGTATCA  
 ACCATCGTCATGGTATTTATTAATACTTCTTCTCCAACAGCTGTTAGCTACTGGGCATATTCATTGATTTCAATCTCTGTA  
 TCAGTAGTTTCAAGTTATGTTATCTGTTCTTACTGTAATAAGATATCTGCACCTAAATATACAACGATTTCATGATATTAC  
 AGAAATTAAGCGGATTCAAGTTGGGAAGTTCGTCACTAA

## 4138.1 (SEQ. ID. NO. 246)

ATGAAATTTAGTAAAAAATATATAGCAGCTGGATCAGCTGTTATCGTATCCTTGAGTCTATGTGCCTATGCACTAAACCAGCA  
 TCGTTTCGACGAAAAATAAGGACAATAATCGTGTCTCTTATGTGGATGGCAGCCAGTCAAGTCAGAAAAAGTGAAAACCTTGACA  
 CCAGACCAGGTTAGCCAGAAAGAAGGAATTCAGGCTGAGCAAAATTGTAATCAAAATACAGATCAGGGCTATGTAACGTCA



CACGGTGACCACTATCATTACTATAATGGGAAAGTTCCTTATGATGCCCTCTTTAGTGAAGAACTCTTGATGAAGGATCCAAA  
 CTATCAACTTAAAGACGCTGATATTGTCAATGAAGTCAAGGGTGGTTATATCATCAAGGTCGATGGAAAAATATTATGTCTAC  
 CTGAAAGATGCAAGCTCATGCTGATAATGTTTCAAGTAAAGATGAAATCAATCGTCAAAAAACAAGAACATGTCAAAGATAAT  
 GAGAAGGTTAACTCTAATGTTGCTGTAGCAAGGTCTCAGGACGATATACGACAAATGATGGTTATGTCTTTAATCCAGCTG  
 ATATTATCGAAGATACGGGTAATGCTTATATCGTTCTCTCATGGAGGTCACTACTACATTCCAAAAGCGATTATCTGTCT  
 AGTGAATTAGCAGCAGCTAAAGCACATCTGGCTGGAAAAAATATGCAACCGAGTCAGTTAAGCTATTCTTCAACAGCTAGTG  
 ACAATAACACGCAATCTGTAGCAAAAGGATCAACTAGCAAGCCAGCAAAATAATCTGAAAACTCCAGAGTCTTTTGAAGG  
 AACTCTATGATTACCTAGCGCCCAACGTTACAGTGAATCAGATGGCCTGGTCTTTGACCCTGCTAAGATTATCAGTCGTACA  
 CCAAATGGAGTTGCGATTCCGCATGGCGACCAATTACCCTTTATTCCTTACAGCAAGCTTTCTGCCTTAGAAGAAAAGATTGC  
 CAGAATGGTGCCTATCAGTGGAACTGGTTCTACAGTTCTACAAATGCAAAACCTAATGAAGTAGTGTCTAGTCTAGGCAGT  
 CTTTCAAGCAATCCTTCTTTTAAACGACAAGTAAGGAGCTCTCTCAGCATCTGATGGTTATATTTTTAATCCAAAAGATATC  
 GTTGAAGAAAACGGCTACAGCTTATATTGTAAGACATGGTGATCATTTCCTTACATTCCAAAATCAAATCAAATTGGGCAAC  
 CGACTCTTCCAAAACAATAGTCTAGCAACACCTTCTCCATCTCTTCCAATCAATCCAGGAACCTTCACATGAGAAACATGAAGA  
 AGATGGATACGGATTGATGCTAATCGTATTATCGCTGAAGATGAATCAGGTTTGTCTAGTACGGAGACCACAATCAT  
 TATTTCTTCAAGAAGGACTTGACAGAAGAGCAAATTAAGGTGCGCAAAAACATTTAG

## 4139.1 (SEQ. ID. NO. 247)

ATGAAAAAAGAGCAATAGTGGCAGTCATTGTACTGCTTTTGAATTGGGCTGGATCAGTTGGTCAAATCCTATATCGTCCAGC  
 AGATTCCACTGGGTGAAGTGGCTCCTGGATCCCAAATTTCTGTTAGCTTGACCTACCTGCAAAATCGAGGTGCAGCCTTTTCT  
 ATCTTACAAGATCAGCAGCTGTTATTCGCTGTCTACTCTGGTTGTCGTGATAGGTGCCATTTGGTATTTACATAAACACAT  
 GGAGGACTCATTCTGGATGGTCTTGGGTTTGACTCTAATAATCGCGGGTGGTCTTGGAACTTTATTGACAGGGTCAGTCAGG  
 GCTTTGTTGTGGATATGTTCCACCTTGACTTTATCAACTTTGCAATTTCAATGTGGCAGATAGCTATCTGACGGTTGGAGTGA  
 TTATTTATTGATTGCAATGCTAAAAGAGGAAATAAATGGAATAA

## 4139.5 (SEQ. ID. NO. 248)

ATGAATACAAATCTTGCAAGTTTTATCGTTGGACTGATCATCGATGAAAACGACCGTTTTTACTTTGTGCAAAAGGATGGTCA  
 AACCTATGCTCTTGCTAAGGAAGAAGGCCAACATACAGTAGGGGATACGGTCAAAGGTTTTGCATACACGGATATGAAGCA  
 AAAACTCCGCCTGACAACCTTAGAAGTGACTGCCACTCAGGACCAATTTGGTTGGGGACGTGTACAGAGGTTCTGTAAGGAC  
 TTGGGTGTCTTTGTGGATACAGGCCTTCTGACAAGGAAATCGTTGTGTCCTCGATATTCTCCCTGAGCTCAAGGAACCTCG  
 GCCTAAGAAGGGCGACCAACTCTACATCCGTCTTGAAGTGGAATAAGAAAGACCGTATCTGGGCGCTCTTGGCTTATCAAGAA  
 GACTTCCAACGTCTTGCTCGTCTGCTACAACAACATGCAGAACCAAACTGGCCAGCCATTGTTTACCGTCTCAAGCTGTC  
 AGGAACCTTTGTTTACCTACCAGAAAAATAATATGCTTGGTTTATTCATCTAGCGAGCGTTACGCAGAGCCACGTTTGGGGC  
 AAGTATTAGATGCGCGCTTATTGGTTTCCGTGAAGTGACCGCACTTGAACCTCTCCCTCAAACCCAGCTCCTTTGAAATG  
 TTGGAACCGATGCTCAGATGATTTTGACTTATTTGAAAGCAATGGCGGTTTCATGACCTTAAATGACAAGTCATCTCCAGA  
 CGACATCAAGGCAACCTTTGGCATTCTAAAGGTCAGTTCAAGAAAGCTTTAGTGGTCTTATGAAGGCTGGTAAAATCAAG  
 CAGGACCAGTTTGGGACAGAGTTGATTTAG

## 4139.8 (SEQ. ID. NO. 249)

ATGAAGATGTTAGTCTATTTTATTGAAAAAAGTTTTCAAAGCCGCTTAAACTGGATTGTCTTAGCTTTATTTGTATCTGTA  
 CTCGGTGTACCTTTTATTTAAATAGTCAGACTGCAAACTCACACGTTGGAGAGCAGGTTGGAAAGTCGCATTGCAGCCA  
 ACGAGAGGGCTATCAATGAAAATGAAGAGAACTCTCCCAAATGTCTGATACCAGCTCGGAGGAAATACCAGTTTGCTAAAA  
 ATAATTTAGACGTGCAAAAAAATCTTTTGACGCGAAAGACAGAAATCTGACTTTATTAAGAAGAGGGCGCTGGAAAGAAG  
 CCTACTATTGTCAGTGGCAAGATGAAGAGAAAGATTATGAATTTGTATCAAAATGACCCGACTGCTAGCCCTGGCTTAAAAAT  
 GGGGTTTGACCGCGAACGGAAGATTACCAAGCCCTGTATCTTGAACATAAAAGCACATACTTTGGGTTTCCGACCCAC  
 GGGATTGATCAGATTGTCTGGATTTAGAGGTTATCATCCCAAGTTTGTGTTGGTTGCTATTATTTTTATGCTAACACAATA  
 TTTGCAAAAGATATCAAAATCATCTGGACACAGCTCACTATATCTGTTTCAAAGTGACATTGCAATATCCTCTCTTTGG  
 AGTTGGAGTGGATATGTAACGTGCTGTTTACGGAATCTGTGGCTTTCTTTCTAGTGGGAAGTCGATAAGTGGTTTGG  
 GACAGTTAGATTATCCCTACCCAATTTATAGCTTAGTGAATCAAGAAGTAACATTTGGGAAAATACAAGTGTATTATTTCT  
 GGCTTGCTCTTAGCTTTCTTAGCCTTATCGTCATTGTGGAAGTTGTGACTTGATTGCTTACTTTTCAAGCAAAAAATGCCT  
 GTCCTCTTTCTTCACTCAATTGGGATTGTTGGCTTATTGTTTGGTATCCAAACCAATTCAGCCTCTTCAAAGGATTGCACATCTG  
 ATTCCTTTACTTTGCTTCAAGTGGAGATTTATCTGGAAGATTACCTAAGCAGATTGATAATGTCGATCTAAATTGGAG  
 CATGGGAATGGTCTTACTTCTTGCCTGATTATCTTTTGTCTATTGGGAATTCTATTTATTGAAAGATGGGGAAGTTTCACAGA  
 AAAAGAATTTTTTAATAGATTCTAG

## 4141.1 (SEQ. ID. NO. 250)

ATGATGAAGTTCATATTGGATATTGTTAGTACACCAGCTATTTTAGTAGCTTTAATTGCAATCTTAGGATTAGTTCTTCAGAA  
 GAAGAAATTACCTGATATTATTAAGGTGGAATTAAGACCTTTGTTGGTTTCTTAGTTGATCTGGTGGTGCAGGAATTGTAC  
 AAAATTCCTTTAAATCCATTGGTACCATTGTTGAGCATGCTTTTCAATTATCTGGCGTTGTGCCGAATAATGAAGCAATTGTAG  
 CTGTAGCTTTAACAACATATGGCTCAGCTACTGCAATGATTATGTTTGCAGGCATGGTGTCAATATCTTAATCGCTCGTTTTA  
 CTCGATTAAATATAATTTTTTAAACAGGGCACCACACTCTATATATGGCATGTATGATTGCGGTCATTTTATCAGTTGCTGGCT  
 TTAAGTCTGCTCTCATCTTACTAGGAGGATTAGCACTCGGTATTATATGAGTATTTCCCCAGCATTTGTGCAAAAAATATA  
 TGGTTCAATTAACCTGGAATGACAAGGTAGCTTTAGGTCAATTCAGTTCTTTGGGATATTGGTTGAGTGGTTTACTGGTAGC  
 CTTATCGGTGACAAATCAAAATCAACAGAGGACATTAATTTCCAAAGAGTTTAGCTTTTTACGTGATAGTACTGTTAGTAT  
 TACTTTATCCATGGCAGTTATTTACATTATTGTAGTATCTTTGCAAGGTCAGAAATATATAGAAAAAGAAATCAGTAGTGGTA  
 CAAGTGGTCTAGTTTATGCTTTTACAATAGCAGGTCAATTTGACAGAGGGGTATTTGTTATTTTAGCAGGTTTTCGCTTATTT  
 TGGGCGAAATTGTTCCAGCCTTTAAAGGTATTTAGAGCGTCTTGTACCTAATTCAAAACCTGCTTTGGATTGTCCGATTGTTT  
 ATACTTATGCAACCAATGCAGTTCTAATTGGATTATCTCTAGTTTGTGGTGGTTAGTAAGTATGGTAATTATGATTGCTT  
 CAGGAACGGTTGTTATCTTACCAGGTGTTGTGCTCAATTTCTCTGTGGAGCGACTGCAGGTGTCATTGGGAATGCATCTGGT  
 GGTGTTCTGGAGCCACTATTGGAGCATTTTACAAGGTATTGTAACAGGTTTCTTCCAGTCTTTTAAATGCCAGTTTGGGA  
 GGACTTGGTTTCCAAGGATCAACTTTCTCAGATGCAGATTTTGGTCTATCAGGAATTATTTAGGAATGTTAAATCAATTTGG  
 CTCACAAGCAGGCATTGTGATTGGTCTTGTCTTATCTAGCAGTTATGTTGGAGTATCCTTTATTAAGAAGCCATCTGCAAC  
 GGAGGAATAA

## 4142.3 (SEQ. ID. NO. 251)

ATGATTA AAAACATTTCTCTCTGCCCTTTCCGGTCATTCTCTTTTCTATCCCTATCATAACTTATTTCTTTTCCCATCTTCTAATCT  
 TAACATTTGGCTATCTACCCAACTATCTTGGCACAGATTTATGCCCTTCCCTTAGCTACTGCAACTATGGCTGCTATTTTAAG  
 TTTCTTATTTTTTTTCTATCTTTTTACAAGAAAAATAAACAAATACGGTTTTACTCTGGCATTTTGGCTTACTATCGCTCATA  
 TTACTATTATTCGGAACAGATAAAACCCCTTTCTCTGCATCAAAATAAGACTAAAACCTTAAAAATTAGTAACCTTGGAAACGTCGC  
 TAATCAAAATAGAAACACAAACATATTGAGCGAATTTTATGCCATTTTGACGCCGATATGGCTATATTTCCCTGAACTAGCTACCA  
 ATATCAGAGGTGAGCAAGAAAACAGAGAATCAAACCTATTGTTTCATCAAGTTGGACTTTCTATGGCCAACTATGATATTTTC  
 ACTTCTCCACCTACCAATAGTGAATAGCTCCTGTGACTGTGATTGTCAAGAAAAAGTTATGGTTTCTATACAGAAGCTAAAAAC  
 TTTTCATACAACACGGTTTCGGGACAATTGTATTACATTGAGAAAAACAAAATATACCAGATATCATTTGCCCTGCATATGCGC  
 CTCCTCTGCCAGGTTTAAATGGAATCTGGAAGCAAGACTTAAACATCATTCATAATCAATTGGCTTCAAAATATCCAAAGGCT  
 ATTATTGCAGGTGATTTTAAATGCAACTATGCGTCATGGAGCACTTGCAAAAAATAAGCTCTCATAGGGACGCATTAAATGCAC  
 TGCCACCTTTTGAAAGAGGAACCTGGAATAGCCAAAGTCCAAAACCTTTTAAATGCAACAATAGATCATATTTTATTGCCTAAA  
 AACCACTACTATGTTAAAGATTAGACATTGTAAGTTTCAAACCTCTGATCATAGATGTATTTTACAGAAAACACATTTTA  
 A

## 4142.4 (SEQ. ID. NO. 252)

ATGAATCCAAATCCAAAGATCTTGGGCTTATGTCAGCAGAAAAGCGACTGAGAAGTTTATTTTATTTCTGATTTTATTGGTCTT  
 ATTGGCCGGAATTTAGCCTGTTTACTCTGATGAAGTCCAACAAAACAGTAGAAAGCAATCTTTATAAATCACTCAATACA  
 TCTTTTTCTATTAAGAAGATAGAGAATGGTCAGACATTCAAGTTGTGACAGCTAGCATCTGTAAGCAAGATTAAGGGGCTGG  
 AAAATGTCTCTCTGAACTTGAGACGGTCGCAAAACTAAAAGACAAGGAAGCACTGGCTGGCGAGCAGCTTTGACGCTG  
 ATGATTTATCAGCTGCAGACAATAACTTGGTTAGCTTAAACGGCTCTTGAGGATTTCATCCAAGGATGTAACCTTTACCAGTTTCG  
 GCTTTCAATCTAAAAGAAGGGCGACACCTTCAAAAAGGGGATTCCAAGAAAAATCCTTATCCACGAAGAATTGGCTAAGAAG  
 AACGGTCTTTTCGCTTCATGACAAGATTGGCTTGGATGCTGGTCAGTCTGAATCTGGAAGGACAAAACAGTAGAGTTTGAGA  
 TTATCGGCATCTTTCTGGTAAAAACAAGAGAAAATTCACAGGCTTGTCTTCTGACTTCACTGGAAGATCAAGCTTTGACGAC  
 TATGAAAGTAGCCAAACCTTTTGGGCAATAGTGAAGCTCAAGTCAGTGCAGCAGCTTCTATGTAGAAAAATCCTAAGGAAA  
 TGGACGGACTCATGAAGCAGGTAGAAAACTTGGCCTTGGAAAATCAAGGCTACCAAGTCGAAAAGGAAAACAAGGCTTTTG  
 AACAAATCAAAGACTAGTTGCAACTTCCAAACCTTCCAGCATTCTCCTTTATGGGATGTTGATAGCAGGAGCTGGAGCC  
 TTAATTCTGGTTTTGTCTCTCTGGTTGAGAGAACGGGTCTATGAAGTGGGGATTTTACTTGCATTGGAAGGCAAGAGCTC  
 GATCTTCTCTACAATTCTGTTTAGAGGTAGTTTGGTATCTCTTGGAGCTTTGCTTCCAGCATTGTTGTCAGGAAACGCAATCAC  
 AACTTACCTACTCCAACTCTACTAGCAAGTGGAGATCAGGCAAGCTTACAAGATACACTAGCCAAAGCAAGCAGTTTATCA  
 ACTAGCATCTTATCTTTTGCAGAATCTATGTTTTCTAGTTCTGCTTAGTTGCTTATCTGTAGCCCTTTGTTTCTTATTCTTAT  
 TAGAAAATCACCGAAAGAAATTTTATCATCTATTAGTTAA

## 4142.5 (SEQ. ID. NO. 253)

ATGTTACACAACGCATTTGCCATGTTACAAGGAAGTTTTCAAATCGATTGTCATCTTCTCTGATTATTCTCCTCATGGCGAGC  
 TTGAGTTTGGTCGGCTTGTCATCAAGGGAGCTACTGCCAAGGCTTCTCAGGAGACCTTTAAAAATATCACCATAGCTTCTC  
 CATGCAAAATCAATCGTCGCTCAACCAAGGAACGCCTCGTGGTCTGGGAATATCAAGGGTGAAGACATCAAAAAATCAC  
 CGAAAACAAGGCCATTGAGTCTTATGTCAAACGTATCAACGCTATCGGAGATTGACTGGATATGACCTGATTGAAACGCCA  
 GAAACCAAGAAGAATCTCACTGCTGATCGTGCCAAGCGTTTGGAAAGTAGCTTGATGATTACAGGTGTCAATGACTCCTCTA  
 AAGAAGACAAGTTTGTCTCTGGTTCTTATAAACTAGTGAAGGAGAGCACTTAAACCAACGACGACAAGGATAAAATCCTCTT  
 GCACAAGGACTTGGCAGCCAAACACGGCTGGAAGTAGGGGACAAGTTAACTGGACTCTAATATCTACGATGCAGATAA  
 TGAATAAGAGGAGCTACAGGAGCAAGTTGAAGTGACAATCAAGGCACTCTTGTAGTGCATATAAAGTACGACGATTAACCT  
 ACAAGAATTTACGAAAACACAGCTATTACAGACATTCACACTGCTGCAAACTTTATGGATACACAGAAGACACAGCCATT  
 TATGGGGACGCAACCTTCTTTGTAACAGCAGACAAGAACTTGGATGATGTTATGAAAGAGTTGAATGGCATCAGTGGTATCA  
 ACTGGAAGAGCTACAGGAGCTGCTCAAGAGCTCCTCTAACTCCAGCTCTTGAGCAATCTATCTCTGGTATGTACAAGATGGCC  
 AACCTCCTCTTCTGGGGTAGCTTGAGCTTCTCAGTTCTCCTCCTTGGCCCTCTTGTCTAGCCTTTGGATCAACGCCCGTCGAAG  
 GAAGTGGGAATTTCTCTCTATCGGCCCTCAAGCAAGCAAGTATCTTGGGTCA  
 ATTCACTACCGAATCTATCTTGATTGCTATCCCTGCTATGTTTCTGCTTACTTCTAGCTAATTACACTGCCCGTGCAATTGG  
 AAACATCTGCTCTTGCATCTTGACTTTCAGGTGTGTTGCCAAACAGGCTAGTAAGGCGGCTCAAGCCTCTAACCTTGGTGGTGGT  
 GCAGAAGTAGATGGCTTTAGCAAGACCTTGTGAGCCTAGACATTTCCATTACAGACATCAGACTTTATCATCATTTTGTCTT  
 TGCCTTGGTTCTAGTGGTTCTCGTTATGGCGCTTGCTTCAAGCAATCTCCTTAGAAAAACAACCAAAAGAGCTCTTGCTGGATG  
 GTGAATAA

## 4144.1 (SEQ. ID. NO. 254)

ATGTCACAGGATAAAACAAATGAAAGCTGTTTCTCCCTTCTGACGAGGTTATCAATATCTCATCGATTGTCGGTGGGGTTGG  
 GAGTTTGATTTTCTGTATTTGGGCTTATCAGGCTGGGATTTTACAATCCAAGGAAACCTCTCTGCCTTTATCCAGCAGGCAG  
 GCATCTGGGGTCCACCTCTCTTTATCTTTTACAGATTTTACAGACTGTCGTCCCTATCATTCCAGGGGCTTGACCTCGGTGG  
 CTGGGGCTCTTATCTACGGGCACATCATCGGGACTATCTACAATATATCGGCATCGTGATTGGCTGTGCCATTATCTTTTATC  
 TAGTGGCCTATACGGAGCTGCCTTTGTCCAGTCTGTCTGACGAAGCGCACCTACGACAAGTACATCGACTGGCTAGATAA  
 GGGCAATCGTTTTGACCGCTTCTTTATTTTATGATGATTTGGCCCATTAGCCAGCTGACTTTCTCTGTATGCTGGCTGCCCT  
 GACCAAGATGAGCTTCAAGCGCTACATGACCATCATCTTCTGACCAACCCCTTACCCTCGTGGTTATACCTACGGTCTGA  
 CCTATATTATTGACTTTTTCTGGCAATGCTTTGA

## 4144.2 (SEQ. ID. NO. 255)

ATGAGAAATATGTTGGTTGTAATCAAGGAAACCTATCTTCGACATGTGAGTCACTGGAGTTTCTTCTTTATGGTGATTTTCGCC  
 GTTCTCTTTTTAGGAATCTCTGTAGGAATTTGGGCATCTCCAAGGTTCTTCTATGGCTAAAAATAATAAAGTGCCAGTAGTGA  
 CAACAGTGCCATCTGTAGCAGAAGGACTGAAGAATGTAAATGGTGTTAACTTCGACTATAAAGACGAAGCAAGTGCCAAAG  
 AAGCAATTAAGAAGAAAAATTAAGGTTATTTGACCATTGATCAAGAAGATAGTGTCTAAAGGCAAGTTATCATGGCGA  
 AACATCGCTTGAATAATGGAATTAATTTGAGGTTACAGGTACACTCAATGAAGTCAAAATCAGCTTAATCGTTCAACTGCT  
 TCCTTGTCTCAAGAGCAGGAAAAACGCTTAGCGCAGACAATTCATTCACAGAAAAAGATTGATGAAGCCAAGGAAAAATAA  
 AAGTTTATTCAAACAATTGCAGCAGGTGCCTTAGGATTCTTTCTTTATATGATTCTGATTACCTATGCGGGTGTAACAGCTCA  
 GGAAGTTGCCAGTGAAAAAGGCACCAAAATTAAGGAAGTCGTTTTCTAGCATAAGGGCAAGTCACTATTCTATGCGCGG  
 ATGATGGCTCTGTTCTAGTATTTAAACGCATATTTGGGATCTATGTTGATGGTGTCTGGCTGCCGTTTGTCTTTAAAGAT  
 TTGCCATTCTTGGCTAGTCTGGTATTTTGGATCACTTGGGAGATGCTATCTCACTGAATACCTTGCTCTTTATTTTATGATCAGT  
 CTTTTCATGTACGTAGCTTGGCAGCCTTCTAGGATCTATGGTTTCTCTGCTGAGGACTCAGGGAAGCCTTGTGCGCTTTG  
 ATGATTTTGATTATGGGTGGTTTTTTTGGAGTGACAGCTTAGGTGCAGCTGGTGACAAATCTCCTCTTGAAGATTGGTTCTTAT



ATTCCCTTTATTTGACCTTCTTTATGCCGTTTCGAACGATTAATGACTATGCGGGGGAGCAGAAGCATGGATTTCACTTGC  
TATTACAGTGATTTTTGCGGTGGTAGCAACAGGATTTATCGGACGCATGTATGCTAGTCTCGTTCTTCAAACGGATGATTTAG  
GGATTTGGAAAAACCTTTAAACGTGCCTTATCTTATAAATAG

4144.3 (SEQ. ID. NO. 256)

ATGACAGAAACCATTAAATTGATGAAGGCTCATACTTCACTGCGCAGGTTTAAAGAGCAAGAAATCCCAAGTAGACTTAA  
ATGAGATTTTGACAGCAGCCAGATGGCATCATCTTGAAGAATTTCCAATCCTACTCTGTGATTGTGGTACGAAGTCAAGA  
GAAGAAAGATGCCTTGTATGAATTGGTACCTCAAGAAGCCATTGCGCAGTCTGCTGTTTTCTCTTTGTGCGGAGATTGA  
ACCGAGCAGAAAAGGGAGCCCGACTTACATACCGACACCTTCCAACCCCAAGGTGTGGAAGGTCTCTTGATTAGTTCGGTCTGA  
TGCAGCTCTTGCTGGACAAAACGCTTGTGGCAGCTGAAAGCTTGGGCTATGGTGGTGTGATTATCGGTTTGGTTCGATACA  
AGTCTGAAGAAGTGGCAGAGCTCTTAACTACCTGACTACACCTATTCTGTCTTTGGGATGGCACTGGGTGTGCCAAATCAA  
CATCATGATATGAAACCGAGAGCTGCCACTAGAGAATGTTGTCTTTGAGGAAGAATACCAAGAACAGTCAACTGAGGCAATCC  
AAGCTTATGACCGTGTTCAGGCTGACTATGCTGGGGCGCTGCGACCACAAGCTGGAGTCAGCGCTAGCAGAACAGTTTGG  
TCAAGCTGAACCAAGCTCAACTAGAAAAAATCTTGAACAGAAGAAATATTATTGATG

4146.1 (SEQ. ID. NO. 257)

ATGTTAAAACTTATTGCTATTGTTGGAACAAATTCAAAACGTTCTACAAAACCGTCAATTGCTTCAATACATGCAAAAAACCTT  
TACTGACAAAGCTGAAATTTGAACCTTGTGAAATCAAGGCCATTCTGTCTTCAACAAACCAGCTGACAAGCAAGTACCTGCT  
GAAATATTGGAAATTGCTGCTAAAAATCGAAGAGGCAGATGGCGTATTATCGGTACTCCTGAGTATGATCACTCTATTCCAG  
CTGTTTTGATGAGCGCTCTTGCTTGGTGTCTTATGGTATTTACCCACTTTTGAACAAACCAATCATGATTACAGGTGCTTCTT  
ACGGTACGCTTGGTTTCATCTCGTGCCCAATTGCAACTTCGTCAAATCTTGAATGCTCCTGAAATCAAGGCAAAATGTTCTTCCA  
GATGAATTTCTGCTCTCACACTCTCTTCAAGCATTTAACCAAGTGGCGACTTGGTTGACCTTGATGTTATCAAGAAATTTGGA  
TGCCATCTTTGATGACTTCCGTATCTTTGTA AAAATCACAGAAAAAATTACGTAATGCACAAGAAATTACTTCGCAAGATGCTG  
AAGACTTTGACTGGGAAAAATTTGTAA

4146.2 (SEQ. ID. NO. 258)

ATGAATACCTATCAATTAATAATGGAGTAGAAAATCCAGTATTGGGATTTGGAACCTTTAAAGGCTAAGGATGGAGAAGAAG  
CCTATCGTGCAGTGTTAGAAAGCCTTGAAGGCTGGTATCGTCATATTGATACGGCGGCGATTATCAGAATGAAGAAAGTGT  
TGGTCAAGCAATCAAGATAGCGGAGTTCACGTGAAGAAATGTTCTGTAATACCAAGCTTTGGAATAGTCAGCAAAACCTAT  
GAGCAAACTCGTCAAGCTTTGGAAAAATCTATAGAAAAACTGGGCTTGGATTATTGGATTGTATTGATTTCATTGGCCGAA  
CCCAAAACCGCTCAGAGAAAATGACGCATGGA AAACTCGCAATGCGGAAGTTTGGAGAGCGATGGAAGACCTCTATCAAGA  
AGGGAAAAATCCGTGCTATCGGCGTACCAATTTTCTTCCCATCATTTGGATGCCTTGCTTGA AACTGCAACTATCGTTCCGT  
CGGTCAATCAAGTTTCGTTGGCGCCAGGTGTGATCAAGATCAAGTCAAGTCTACTGTCGTGAAAAGGGAATTTTATTGGA  
AGCTTGGGGGCTTTTGGACAAAGGAGAACTGTTTGTAGCAAGCAAGTCCAAGAAATAGCAGCAAAATCAGGAAAAATCGGT  
TGCTCAGATAGCCTTGGCCTGGAGCTTGGCAGAAAGGATTTTACCACCTTCCAAAATCTGTCACAACCTCTCGTATTTCAAGCTA  
ATCTTGATTGCTTTGGAATTGAACGTAGTTCATGAGGAGAGAGAAACCTTAAAAACGATTGCTGTTCAATCGGGTGCTCCACG  
AGTTGATGATGTGGATTTCTAG

4147.1 (SEQ. ID. NO. 259)

ATGAGGTGCAAAATCTTGATCCAATTGCTATTCAACTAGGACCCCTAGCCATTCTGTTGGTATGCCTTATGTATTGTGACAGG  
CTTGATCTTGGGTTTATTGACCATGAAAGAAAGCACCTAGAAAGAAGATCATACCAGACGATATTTAGATTTTATCTTAG  
TAGCCTTTCCCTTGGCTATTTTAGGAGCTCGTCTCTACTATGTTATTTCCGATTGATTACTATAGTCAGAAATTTAGGAGAGA  
TTTTTGCCATTTGGAATGGTGGTTTGGCCATTTACGGTGGTTTGAATACTGGGGCTCTTGCTGCTCTATATCTTTGCTGACCGTA  
AACTCATCAATACTTTGGGATTTTCTAGATATTGCGGCGCCTAGCGTTATGATTGCTCAAAGTTTGGGGCGTTGGGGTAATTTT  
TTTAAACCAAGAAGCTTATGGTGCAACAGTGGATAATCTGGATTATCTACCTGGCTTTATCCGTGACCAGATGTATATTGAGGG  
GAGCTACCGTCAACCGACTTTCTTTATGAGTCTCTATGGAATCTGCTTGGCTTTGCCTTGATTCTGATTTTAGACGGAAATG  
GAAGAGTCTCAGACGAGGTCAATACACGGCCTTTACTTGATTGTTGGTATGGTTTTCGGTTCGATGGTTATCGAAGGTATGCGAA  
CAGATAGTCTCATGTTCTTCGGCTTTGAGTGTCCCAATGGCTTTCAGTTGCTCTTATCGGTCTCGGTATAATGATCGTTATTT  
ATCAAAATCGAAAGAAGGCCCTTACTATATTACAGAGGAGGAAAACTAA

4147.2 (SEQ. ID. NO. 260)

ATGGGTAAATTATCCTCAATCCTTTTAGGAACCGTTTCAGGTGCAGCTCTTGCTTGTTTTTAACAAGTGATAAGGGCAAACA  
AGTTTGCAAGTCAAGCTCAAGATTTTCTAGATGATTGAGAGAAGATCCGGAGTATGCCAAGGAGCAAGTCTGTGAAAAACTG  
ACAGAAGTTAAGGAGCAGGCTACAGATTTTGTCTGAAAACAAAAGAACAGGTTGAGTCAGGTGAAATCACTGTGGACAGT  
ATACTTGCTCAAACTAAATCCTATGCTTTTCAAGCGACAGAAGCATCAAAAAATCAATTAATAATCTCAAGGAGCAATGGC  
AAGAAAAAGCCGAAGCTCTTGATGACTCAGAAGAGATTGTGATTGATATAACAGAAGAATAA

4147.3 (SEQ. ID. NO. 261)

ATGAAAACATAAATTGATCTTTTGGGGCTCTATGCTCTTTCTCTCTCCCTCTCCATCCTTCTGACCATTTATCTGGCTTGGATTT  
TCTATCCTATGGAGATTCACTGGCTAAACCTTAACGAATCGAGTCTATCTAAAACCAAGAAACCAATCAATACAATTTTCATATC  
TTGATGAATTATCTGACCAATCCTTTTATGTCAGGTCTTACAGATGCCTGATTTTCGTTTCGTCAGCAGCTGGTCTGCACCATTT  
GCAGTGGTCAAGAATCTCTTTCATTTGGTTTCAGCTAGTAGTCTAGTGACACTGCCAAGTTTCTATGCTTTTGTCAATAGGATT  
GTGAAAAAGGACTTTTGTCTCTTTATCGAAAAAGTCTCCTGGCTCTAGTAGTCTTACCTGTGATGATTGGACTTGGGGGAGT  
TTTGATTGGTTTGGACCAATCTTACTCTTTTCCATCAAAATCTCTTTTGTGGGAGATGATACCTGGCTTTTGTATCCAGCCAA  
GGATCCTGTTATTGATTTTGGCAGAGACTTCTTTCTCATGCCTCTCTCTTTTGTCCCTCTATGAAAACCTCTTTGGC  
TATCTGTATCTGAAAAGTCGTAGGAAGTGA

4149.1 (SEQ. ID. NO. 262)

ATGACTTATCATTTTACTGAAGAATACGATATTATTGTAATTGGTGCGGGACACGCTGGGGTTGAGGCTTCCTTGGCCGCTAG  
CCGTATGGGCTGTAAGGTCTGCTTGGCACCATCAATATTGAAATGCTGGCTTTTCATGCCTTGTAAATCCCTCTATCGGTGGTTC  
TGCCAAGGGGATTGTCGTGCGTGAAGTGCATGCCCTCGTGGCGAGATGGCCAAAACCAATGACAAGACTTACATCCAGATG  
AAGATGCTAAACACAGGGAAGGGCCAGCTGTCCGTGCCCTTCGTGCGCAGGCTGACAAGGAACCTTACTCTAAGGAGATG  
CGCAAGACGGTTGAAAAACCAAGAAATCTGACCCCTTCGTCAAACCATGATTGATGAGATTTTGGTGGAAAGATGGCAAGGTTG  
TCGGTGTGCGTACAGCCACCAATCAAGAATATGCTGCTAAGGCTGTTATTGTGACGACAGGAGTCTCTCCGTGGGGAAAT  
TATCATCGGAGACCTCAAGTACTCATCAGGTCTTAACACAGCTTGGCTTCTATTAACCTAGCTGACAATCTCAAGGAAGTGG

GTCTCGAAATCGGTGCTTTCAAGACAGGAACCCCTCCACGTGTCAAGGCTTCTTCTATCAATTACGATGTGACAGAAATTCAG  
 CCAGGAGACGAAGTGCCTAATCATTTCTCATACACTTCACGTGATGAGGATTATGTCAAGGACCAAGTACCATGCTGGTTGA  
 CCTATACCAATGGTACCAGTCATGAGATTATCCAAAACAACCTCCACCGTGCCTATGTTTACAGGTTGGTCAAGGGAAT  
 GGGGCTCGTTACTGTCCGTGATTGAAGACAAGATTGTGCGCTTTGCGGACAAGGAACGTACCAACTCTTCTTCTGAGCCA  
 GAAGGGCGCAACTACTGAGGAAGTCTATGTGCAAGGACTTTCAACCAAGTCTGCCTGAGGATGTCCAGCGTACTTGGTGATT  
 CCATCAAAGGTTTGGAAATGCAGAGATGATGCGGACAGGTTATGCTATTGA  
 GTATGATATGGTCTTGCCTCATCAGTTGCGTGCGACTTTGGAAACCAAGAAAAATCTCAGGTCTCTTCACTGCTGGTCAGACAA  
 ATGGAACATCAGGTTACGAAGAGGCAGCAGGCCAAGGATTATCGCGGGTATCAATGCGGCTCTGAAAAATCCAAGGCAAGC  
 CTGAATTGATTTTGAAGCGCAGTGATGGTTATATCGGGGTGATGATCGACGACTTGGTGACCAAGGGAACCATTTGAACCTTA  
 CCGTCTCTTGACCAGTCGTGCTGAATACCGTCTCATTTCTCGTCATGACAATGCTGATATGCGCTTGACTGAGATGGGACGCG  
 AGATTGGCCTTGTGGACGATGAACGCTGGGCTCGTTTTGAAATCAAGAAAAATCAATTTGATAATGAGATGAAGCGCCTAGA  
 CAGTATCAAACCTCAAGCCAGTCAAGGAAACCAATGCCAAGGTTGAGGAGATGGGCTTCAAACCTTGACCGATGCGAGTAC  
 AGCCAAGGAATTCCTTCGCCGTCCAGAAGTTTCTTACCAAGATGTGGTGGCCTTCATCGGACCAGCTGCAGAAGACTTGGAT  
 GACAAGATTATCGAATTGATTGAACAGAAATCAAGTATGAAGGCTATATTTCCAAAGCCATGGACCAGGTTGCCAAGATGA  
 AACGCATGGAAGAAAAACGCATTCCGGCCAATATCGACTGGGATGACATTGATTCTATCGCAACCGAAGCCGTCAGAAGTT  
 CAAACTCATCAAGAAACCATCGGCCAAGCCAGCGTATTTGCGAGTAACCCAGCAGATATTTCTATTTTGTATGGTG  
 TATCTGGAAGGTAAAAATCGTAGTATTTCTAAAACCTTCAAAAATCAAAATGA

## 4149.2 (SEQ. ID. NO. 263)

ATGAAAGTATTAGCTTTTGATACGTCCAGCAAGGCTCTTTCTCTGGCTATTTTAGAGGATAAGCAGGTTCTTGCCGAGACGAC  
 GATTAATATTAAGAAAAATCACAGTATTACTCTTATGCTGCCATCGATTTTTTGATGGCAAGTTTGGATTGGACACCCAAGG  
 ATTTGGACCGAATCGTGGTAGCTGAAGGGCCGGGTAGCTATACAGGCTTGCGAATTGCGGTAGCAACTGCTAAGACCTTAGC  
 TCACACCTTGAACATCGAGTTGGTTGGTATGTGCGAGTCTCTTGGCTCTGGTGCCCATCAACAAGAAGGTTTGTGTGCCCT  
 TGATGGATGCGCGTCGCAATAATGTTTATGCAAGGATTTATGAAAATGCCAAACCTGTCATGGCAGAAGCGCACCTATCTTTT  
 GAAGAGGTGCTAGAAAAAGTCAAGGGTACTAGTCAGGTAACCTTTGTCGGAGAAGTTGGCCCTTTGTTGAGCAGATTCAAA  
 AACACTTGCCAAGGACTGATTACAAGAAACATTGCCAATGCGCTAATCTAGCTCTTTTGGCCTGGGACAAGGAAGCAGA  
 CTCCTTGCATGATTTTGTGCCGAATTACCTCAAACGAGTCGAGGCTGAGGAAAACTGGCTCAAGAACCATACCGAGTCTGGC  
 GAGTCTTACATTAACGCCTATGA

## 4149.3 (SEQ. ID. NO. 264)

ATGATAGAAATCAAGCGAATTCAACAACAGCCTGACCTAGCTCAAGCCATCTACGCTGTTATGGCAGCTGTTTACCTAGTCA  
 GTCTTGGACTCTGGAGCAAAATCCAAGCAGATCTGTCCCAAGACCAGACTTGGTATGCATTGGCTTATGATGGGGCAGAAGT  
 GATTGGATTCTAGCTGTGCAAGGAGAATCTTTTGAAGCAGAAGTCTGCAAAATCGCTGTCAAAGGAGCTTATCAGGGTCAG  
 GGGATTGCGTCagCCTTGTGTTGCTCAATTGCCGACAGACAAGGAAATTTTCTCGAAGTCAGACAGTCAAATCAACGAGCGCA  
 AGCATTTTACAAGAAAGAAAAAGATGACAGTTATCGCTGAGCGAAAGGCCACTACCATGACCCAGTCGAGGACGCCATTATC  
 ATGAAGAGAGAAATAGATGAAGGATAG

## 4152.2 (SEQ. ID. NO. 265)

ATGACAAAACAAGTCTTATTAGTGGATGATGAAGAACACATTCTGAAATTGCTTGACTACCATTTAAGTAAGGAAGGCTTTT  
 CTACTCAATTGGTGACAAATGGACGGAAGGCCCTTAGCTTTGGCAGAAACAGAACCCCTTGATTTTATCTTGCTTGATATCATG  
 TTACCACAATTAGATGGCATGGAAGTTTGTAAAGCGGCTGAGAGCCAAAGGCGTCAAACTCCAATTATGATGGTTTCTGCGA  
 AAAGTATGAAATTTGATAAGGTTTGGCCTTGGAATTAGGGGCTGATGACTACCTGACCAAGCCCTTTAGCCCTAGAGAATT  
 GCTGGCGCGTGTCAAGGCTGTCTCAGGCGAACTAAAGGAGAACAAAGGAGATGATTAGATAATATCGCTGACGATTC  
 TTGGCTATTTGGGACCTTGAAAGTATACCCTGAGCGTCATGAAGTCTACAAGGCGAATAAGTTACTGAGTTTGACCCCAAAA  
 GAATTTGAAAGCGATAAAAAATCCGTTTTTTGAAGTTTTCAAAGTTTCGAAAGTAACCGCCCAATAA

## 4154.1 (SEQ. ID. NO. 266)

ATGACTACTTTTAAAGATGGATTTTTATGGGGTGGTGTCTGTTGCTGCTCATCAACTTGAAGGTGGATGGCAAGAAGGTGGCA  
 AGGGAATTAGTGTGCTGATGTTATGACTGCTGGTGTGTCATGGAGTAGCTCGTGAAATTACTTTGGGAGTTTTAGAGGGTAAA  
 TATTATCCAAATCATGAGGCGATAGATTTTTATCACCGTTATAAAGAAGATATAGCACTTTTTGCTGAAATGGGATTCAAGTG  
 CTTCCGTACCTCTATTGTCATGGACACGTATCTTTCCAAAAGGTGATGAGTTAGAGCCGAATGAAGAAGGATTACAGTTTTATG  
 ATAATCTTTTGTGATGAATGCTTAAAGAATGGTATTGAACCTGTCATCACTCTATCTCATTTTGAATGCCTTATCACTTAGTGA  
 CCGAATATGGTGGTTGGAAAAATAGGAAATTGATTGATTCTTTGCTCGTTTTGCGAGAAGTCGTATTTAAACGTTACAAAGAT  
 AAGGTTAAATATTGGATGACTTTCAATGAAATCAATAATCAAGCGAATTATCAGGAAGATTTTGACCATTTACTAATCACTAG  
 GTATTGTATATGAGGAAGGTGATAATAGAGAAGCAATTATGTATCAAGCAGCACATTACGAATTAGTTGCTTCTGCACGAGC  
 TGTAAAAATTGGTCATGAGATTAATCCAGATTTTCAAATAGGTTGATGATTGCGATGTGTCGAATTTATCCAGTTACTTGCA  
 ATCCTAAGGATATCTTAATGGCAATGAAAGCTATGCAGAAGCGTTATTATTTTGTGATGTGTCATGTTTTAGGTAATATCCT  
 GAGCATATTTTCAAGTATTGGGAACGAAAAGGTATTTCAAGTTGATTTTACTGCCAGGATAAAGAAGATTTACTTGGTGGGA  
 CTGTAGATTACATTGGTTTCACTTACTATATGTCCTTTGCTATCGACTCTCATCGTGAAAATAATCCTTATTTGATTATCTTG  
 AAACAGAAGATTAGTGAAAAATAATTATGTTAAGGCTTCTGAATGGGAGTGGCAAAATTGATCCAGAAGGTTTGGCTTATGC  
 GTTAAATTGGTTTACAGACCACTATCACTTACCCTCTTTATTGTTGAAAATGGTTTGGAGCTATAGATCAAGTTGCAGCAG  
 ATGGTATGGTACATGATGATTATAGAATTGAATATCTAGGTGCCATATCTCGTGAAATGAAAAAGGCTGTAGTTGAAGATGG  
 TGTGATTAAATGGGTTATATCCATGGGGATGATTGATTGGTTTTCAGTGTGACCGGTGAATGCGGAAACGTTATGGCT  
 TTATTTATGTAGATAAAGATGATAATGGGAAGGGAAGTTATAATCGTTCCCCGAAAAAATCTTTTGGCTGGTATAAGGAAGT  
 TATTTTCATCTAACGGTGAATCAGTAGAATAG

## 4154.2 (SEQ. ID. NO. 267)

ATGGATCAACAAAACGGGTTGTTTGGTTTTCTTGAAAACCATGTTATGGGACCAATGGGCAAACTTGCTCAGTTTAAAGTAGT  
 ACGTGCTATCACGGCTGCAGGTATGGCTGCTGTACCATTTACTATTGTAGGATCAATGTTTTGGTATTCAGTATTTGCCACA  
 AGCTTTCTCATTTTGGCCAAATGTGGCAGATATTTTCTGCTTCATTTGATAAATTCACATCACTTTACATGGTTGCCAACTA  
 TGCGCATATGGGTTCTCTATCTCTTATTTTCGTTCTATCACTTGCATATGAATTGACAAAAATTTATGACAGAGGAAGAAGAAC  
 TCAATATGAATCCTCTTAATGGTGCTTGTGCTTGTATGGCTTTTGTATGACAGTACCGCAAAATCATTTTGTATGGTGGAA  
 TGATGAAGACTGTGACAAGTCTAAAAAGAGGTGCAGTAATTGCAGATGGATGGGCAATGGGAAATGTAGTCGCACGTTTTG



GGACAACAGGGATTTTACCGCAATCATTATGGCAATTGTGACTGTTCTATTTATCGTATGTGTGTTAAACATAATTGGGTT  
ATTAATAATGCCTGAAGCTGTTCCAGAAGGAGTTTCTCGTGGAATTTACCGCTTTGGTTCCGGGATTTGTTGTTGCATTTGTTGTT  
ATCTTTATCAACGGTCTTCTGTAGCAATGGGAACAGATATTTTAAAGTCATTGCAATTCCATTTGGTTTGTATCCAATCTG  
ACTAATTCGTGGATTGGTTAATGATTATTTATCTATTGACTCAACTACTTTGGATTGTAGGTATCCACGGTGCGAACATTGTT  
TTTGATTTGTTAGTCCAATTGCTCTTGCTAACATGGCTGAAATGCTGCTGGCGGGCACTTCGCTGTTGCAAGGTGAATTTCT  
AATATGTTTGTAATTGCAGGTGGTCTGGTGCAACTTTAGGACTATGTTTATATATTGCTTTTGCCTCTAAATCTGAACAGCTT  
AAAGCAATAGGACGAGCATCTGTAGTTCACGCCCTTATTTAATATTAATGAACCATTAAATTTTGGATTACCTATTATCTATAA  
TCCAGCCTTGGCTATACCATTATTTAGCACCAATGGTACTGCTACTATTTATTACGTAGCGAATCTCTAAACTTTATTAA  
GCCAATTATCCGACAGGTTCCATGGCCAACTCCAGTAGGGATTTGGAGCTTCTTAGGGACAGCAGATCTTCGAGCTGTATTA  
GTTGCTCTAGTATGTGCATTTGCAGCATCTCTAGTCTATCTCCATTATCCGTGTATATGATCAAAAATTGGTGAAAGAAGA  
GCAAGGTATCTAA

4155.1 (SEQ. ID. NO. 268)

ATGAAAAAATTTTATGTAAGTCCAATTTTCTTATTCTAGTAGGATTGATTGCGTTTGGAGTCTTATCCACTTTCATTATTTTT  
GTTAATAATAATCTGTTGACGGTTTAAATTTGTTTCTTTTGTAGGAGGCTATGTTTTTTTATTTAAGAACTTGAGAGTGCAT  
TATACAAGGAGTGATGTAGAACAGATACAGTATCTAAACCACCAAGCGGAAGAAAGTTTGACAGCTATTGGACACAGATG  
CCTGTAGGTGTTATGAAATTGAATTTATCTTCTGGAGAGGTTGAGTGGTTTAAATCCCTATGCTGAATTGATTTTGACCAAGGA  
AGATGGTGATTTGATTTAGAAGCTGTTCAAACGATTATCAAGGCTTCAGTAGGAAATCCGCTCTACTTATGCCAAGCTTGGTG  
AGAAGCGTTATGCTGTTTATGGAATGCTTCTCCGGTGTTTGTATTTTGTAGATGTATCCAGGGAACAAGCCATAACAGAT  
GAATTGGTAACAAGTAGACCAGTGATTGGGATTGTCTCTGTGGATAATTATGATGATTTGGAGGATGAAACTTCTGAGTCAG  
ATATTAGTCAAAATCAATAGTTTGTAGCTAATTTATATCAGAGTTTTCAGAAAAACACATGATGTTTTCTCGTCGGGTAAGT  
ATGGATCGATTTTATCTATTTACTGACTACACGGTGCTTGAGGGCTTGATGAATGATAAATTTCTGTTATTGATGCTTTCAGA  
GAAGATCGAAACAGAGACAGTTGCCCTTGACCTTAAAGTATGGGATTTTCTTATGGCGATGGAAATCATGATGAGATAGGGA  
AAGTTGCTTTGCTCAATTTGAACCTTGGCTGAAGTACGTGGTGGCGACCAGGTGGTTGTTAAGGAAAACGACGAAACGAAAAA  
TCCAGTTTATTTTGGTGGTGGGCTGCTGCTTCAATCAAGCGTACACGGACTCGTACGCCGCTATGATGACAGCTATTTTCAG  
ATAAGATTCCGAGTGATAGTCAAGTTTGTGATGCTCGTCAAAAAATTTAGACATGGATGCTTTGGGCTTGTCTGATGATTG  
CAGTTGTTGCCAGCAATGTGATTGAAAAATAGCTATGCTCTTTATGATGAAGAACAAATGTCTCCAGATTTGAAACGAGCTGT  
TTCATTCATAGAAAAAGAAGGAGTTACGAAGTTGTTGTCTGTTAAGGATGCAATGGGGATGGTGACCAATCGTCTTTGTTG  
ATTCCTTGTAGACCAATTCAAAGACAGCCTTAACATTATCAAAAGAAATTTTATGATTTATTTACCCAAACCAATGTTATTGACCA  
CCATAGAAGGGATCAGGATTTTCCAGATAATGCGTTTATTCTATATCGAAAGTGGTGCAAGTAGTCCAGTGAGTTGGTA  
ACGGAATTGATTACAGTTCAGAAATCTAAGAAAAATCGTTTGAAGTCGATGCAAGCAAGTGTCTTGTATGGCTGGTATGATGTT  
GGATACTAAAAATTTACCTCGCGAGTAACATAGTCGGACATTTGATGTTGCTAGCTATCTCAGAACCGCGCGGAAGTGATAGT  
ATTGCTATCCAGGAAATCGCTGCGACAGATTTGAAGAATATCGTGAGGTCAATGAACCTATTTTACAGGGGCGTAAATTAG  
GTTCAGATGTACTAATAGCAGAGGCTAAGGACATGAAATGCTATGATACAGTTGTTATTAGTAAGGCAGCAGATGCCATGTT  
AGCCATGTCAGGTATTGAAGCGAGTTTGTCTTGGCAAGAATACACAAGGATTATCTCTATCTCAGCTCGAAGTCGTAGTA  
AACTGAATGTACAACGGATTATGGAAGAGTTAGGCGGTGGAGGCCACTTTAATTTGGCAGCAGCTCAAATTAAGATGTAAC  
CTTGTCAGAAGCAGGTGAAAAACTGACAGAAATGTATTAATGAAATGAAGGAAAAGGAGAAAGAAGATGA

4156.1 (SEQ. ID. NO. 269)

ATGAAAGAGAAAAATATGTGGAAAGAATTGTTGAATCGTGCAGGCTGGATTTTGGTCTTTTACTTGCCGCTCTTTTATATCA  
GGTTCCTTACGTGGTTACCTCTATTTTGACTTTAAAGAAAGTAGCCCTGCTACAGTCAGGGCTGATAGTTGCTGGCCTTTCAA  
TTGTGGTTCTGGCTCTATTTATTATGGGAGCTCGTAAAACCAAGTTAGCTAGTTTAAATTTTCTTTTATAGAGCTAAAGATT  
TGGCAGCTTTGGGCTTGAGTTATCTAGTTATTTGCTGGGTCAAATATCTTGGTTCCATTTTATGCAACTGTCAAATGAGACG  
ACAACAGCTAACCAAGTCAAGATTAAATGATATGGTTCAAAAAATAGTTCGTTGATTTCAGTTTCTTCTGTGCTAGCCTTGTGCT  
CCGATTTGTGAGGAAATCTTGTGCTGTTGGGATTGTTTCTTAAAGATTTCAGAGGCAAGGAGAACTTGGGATTTGTAGTCG  
GTACGATTGTGTTTGCTTTATTGTCATCAACCAAGTAATTTACCTTCTTTATGATTTATGGAGGTATGTGACAGTTCTATCTT  
GGACAGCCTACAAGACCCAACGTTTGGAAATGTGATCTTGGTTTACATGATTGTTAATGGGATTGCTTCTGTTTGTGGCT  
CTTGTTGGTGAATTATGAGTCGGACATTAGGAATTTCTGTTTAAATGAAAGAGAAAAATATGTGGAAGAAGATTGTTGAATCGTG  
CAGGCTGGATTTTGGTCTTTTACTTGCCGCTCTTTTATATCAGGTTCCCTAGTGGTTACCTCTATTTTGACTTTAAAGAAAG  
TAGCCCTGTACAGTCAAGGCTGATAGTTGCTGGCTTTCAATTTGTTGTTCTGGCTCTATTTATTTAGGAGTCTGTAATAAAC  
AAGTTAGCTAGTTTAAATTTTCTTTTATAGAGCTAAAGATTGGCAGCTTTGGGCTTGAGTTATCTAGTTATGCTGGGTCA  
AATATACTTGGTTCCATTTTATTGCAACTGTCAAATGAGACGACAACAGCTAACCAGTCTCAGATTAATGATATGGTTCAAAA  
TAGTTCTGTGATTTCAGTTTCTTCTGCTAGCCTTGTGCTCCGATTTGTGAGGAAATCTTGTGCTGGGATTGTTCTCTAA  
AAAGATTTTCCGAGGCAAGGAGAACTTGGGATTTGTAGTCGGTACGATTGTGTTTGTCTTATTGTCATCAACCAAGTAATTTAC  
CTTCTTATTGATTATGAGGATATGTCGACAGTTCTATCTTGGACAGCCTACAAGACCCCAACGTTTGGAAATGTGATCTTCTG  
CTTCACATGATTGTTAATGGGATTGCTTCTGTTTGTGGCTCTTGTGGTGATTATGAGTCGGACATTAGGAATTTCTGTTTAA

4156.4 (SEQ. ID. NO. 270)

ATGGATACACAAAAGATTGAAGCGGCTGTAAAAATGATTATCGAGGCTGTAGGAGAGGACGCTAATCGCGAGGGCTTGCAG  
GAAACACCTGCTCGTGTAGCCCGTATGTATCAAGAGATTTTTCAGGCTTGGTCAAACAGCAGAGGAACATTTGTCAAAAT  
CCTTTGAAATATTGACGATAATATGGTGGTAGAAAAGGATATCTTTTCCATACCATGTGTGAACACCACTTCTTGCCATTTT  
ATGGTAGAGCGCACATTGCCATACATCCAGATGGTCTGTGGCAGGCTTGCTAAGCTAGCCCGTACGTTTGAAGTTTATTCG  
AAAAAACCAACAAATTCAGAAACGTTTGAATATCGAAGTGGCCGATGCCTTGATGGACTATCTAGGTGCTAAAGGAGCCTTGT  
TTGTCATTGAGGCGGAACATATGTGTATGAGTATGCGTGGTGTAGAAAACCAAGGCACTGCAACCTTGACGACAGTAGCTCG  
TGGTCTATTTGAAACAGATAAGGATCTCCGTGACCAAGCTTATCGTTTAAATGGGGCTATAA

4157.2 (SEQ. ID. NO. 271)

ATGAAAGACTTGTTTTAAAGAGAAAGCAGGCCTTTTCGTAAGGAGTGTCTTGGTTATCTGCGCTATGTGCTCAATGACCACTT  
TGCTTGTCTGCTTGTCTGTTGGGCTTTCTAGCCTACCAGTACAGTCAACTCTTACAACATTTTCCTGAAAAATCATTGGCC  
TATCCTTTTGTGTAGGAATTACGCTGTTTACTTTTACTTTGGGGAGGAACTGCCACCTATATGGAGGCTCCAGACAAGCT  
CTTCTCTTAGTTGGAGAAGAGGAAATTAAGCTCCATCTCAAGCGTCAAACTGGCATTTCCTAGTCTTTTGGCTCTTTGTACA  
GACCTTTTCTTGTGTTATTTGCGCCTTATTTTAGCAATGGGTTATGGCTTGGCAGTTTCTGCTCTATGTGCTTTTATTG  
GGGCTAGGAAATTTTCCACTTTTGTCAAAGGCCAGCAAAATTTTCACTGAAACTGGAGTGGACTGGACTGTATTTATTTC  
TCAAGAAAGCAAGCGTAAGCAAGTCTTGTCTGTTTCTTGGCCTTTTACGCAGGTCAAGGGAATTTCAAACAGCGTTAAGC  
GTCGTGCTATCTGGACTTTATTTTAAAGGCTGTTTCAAGAGTGCTGGGAAGATTGGCAAAATCTCTATCTGCGTCTTAT

CTGCGAAATGGCGACCTCTTTGCTCTCAGTCTTCGTCTTCTCTTCTGCTTTCTTCTGCTGGCGCAGGTTTTTATCGAGCAAGCTTGG  
ATTGCGACAGCAGTGGTAGTCTCTTTAACTACCTCTTGCTCTTCCAGTTGCTGGCCCTCTATCATGCCTTTGACTACCAGTAT  
TTGACCCAACTCTTTCCGCTGGACAAGGGGCAAAAGGAAAAAGGCTTACAGGAGGTAGTTCGAGGATTGACCAGTTTTGTTT  
TACTTGTGGAATTAGTTGTTGGGTTGATTACCTTCCAAGAAAACTAGCCCTTCTAGCCTTACTAGGAGCTGGTTTGGTTTTAC  
TAGTCTTGATTTTGCCTTATCAGGTAAAACGTACAGTGCAGGACTAA

## 4158.1 (SEQ. ID. NO. 272)

ATGAGAAAAATCAATAGATTATTAGCGGCAGATAATGCCTATCTTATTCTTTAGAGACGACTATAAAGTCTGTATTGTATCACAA  
TAGAGATGTTGATTTTATATTCTCAACAGTGATATAGCTCCTGAATGGTTTAAATTTATTGGGGAGAAAAATGGAAGTTGTGA  
ATTCTACAATTCGCAGTGTACACATTGATAAAGAAGCTTTTGAAGCTATAAAACAGGACCTCATATAAATTATGCTTCTTAC  
TTTAGATTTTTTGGCAGACAGAAGTGGTTGAATCTGATAGGGTATTGTATCTGGATTCCGATATCATTGTAAGTGGGAACTAGC  
TACTTTGTTTGAGATAGATCTCAAAGGATATTCAATTTGGTGCTGTTGATGATGTCTATGCCTATGAAGGACGAAAAATCTGGAT  
TTAATACTGGTATGTTACTAATGGATGTTGCAAGTGGAAAGAACATTCTATTGTCAATAGTTTATTGGAATTAGCGGCCGAG  
CAGAATCAAGTTGTTTCATCTTGGGGATCAGAGTATTTAAATATTTATTTTGAGGATAAATTGGCTAGCCTTAGATAAAACATA  
TAATTTATATGGTGGGTATTGATATTTATCACCTTGCTCAAGAATGTGAACGTCTAGATGACAATCCACCTACAATTGTTTCACT  
ATGCTATAGCAGGAACCTTGGAAATACATATAGTATATCTAGACTAGCTGAATTATGGTGGGTTTATAGAGATTGGATTGG  
TCAGAGATTGCTTTTCAACGTTCCGATTTAAATATTTTGAAGAAAGCAATCAGTCTAAAAACAAGTGATGCTTGTGACATG  
GAGTGCAGATATAAAACATTTAGAGTATTTAGTACAACGGTTACCTGATTGGCATTTTCATTGCTGCACCGTGTGATTGTT  
CTGAGGAGCTGACCTCTCTATCAGTATACGAATGTAAACAGTATATCAAAATGTATTACATAGTAGAATTGATTGGCTATTG  
GACGATTCTATAGTTTATTTAGATATTAATACAGGTGGAGAGGTTTTTAATGTAGTTACAAGGGCACAAGAAAGTGGCAAGA  
AAATCTTCCGCTTTTGATATCACACGTAAGATGATGGATGATGGACTCTATGACGGTATTTTTTCTGTGGAGAGACCAGATGAT  
TTAGTGGATAGAATGAAGAATATAGAGATAGAGTAA

## 4158.2 (SEQ. ID. NO. 273)

ATGACTAAGATTTATTCGTCATAGCAGTAAAAAAGGACTATTTACCTCATTTCTACTGTTTATCTATGTATTGGGAAGTCG  
TATTTACTCTCCCTTTGTTGACCTAAATACTAAAGATTTTTAGGAGGTTCAACAGCCTATCTAGCCTTCTAGCCGCCCTAAC  
AGGTGGGAATCTAAGAAAGTTTATCAATTTTCTGTTGGATTATCCCTTGGATGTCGCCCATGATTTTATGGCAGATGTTTTCT  
TTTTTCTAAACGGTTGGGTTTAAACATCTACGCTATAGAAATACAAGATCGCCGTAAAAATGTACCTGACCTTGCTAATTGCTG  
TGATTCAATCCTTGGCAGTTAGCTTGAGACTGCCAGTACAATCCTCTCTATTCTGCAATATTGGTTGTTCTAATGAATACAATAT  
TGCTGATAGCAGGAACATTTTCTGTTTGGTTGTCAGATTAAATGCGAGTATGGGGATTGGAGGTTCTATTGTGAATCCTCC  
TATCCAGTATGGTTTTAAATATTCTCAGGATGTTTTGGAACATTTTCAGACAGTACACATTCCAACAGGGATTATTGTGTTA  
CTTGCTTTATTAACCTTGTCTTTTCTTATTACTTGCCCTTATGTATCGAGCTCGCTATTGGTTCTGTTAAATAAATTTGGCT  
TACACAATCGATTTAAACGCTATTCTTATCTCGAAATCATGTTGAATCCTGCAGGTGGGATGCCTTATATGTATGTATGAGT  
TTTTTAGTGACCAAGCTTATTGTTTCATCTTGTGGGATTATTTTCCCTAATCATTGAGGTTAGCGGCTTTATGAAAGGAA  
TTTATGGTTGGAAGCCTTTGTGGGTCTATGTTTATATTCGGTCTTATTTTATTAGTATCATTTTGTCTTTGTTTACGATGA  
ATGGAGAAGAGATTGCGACCGTATGAAAAATCTGGAGAATACATTTATGGTATTTATCCAGGTGCGGATAGTACGATT  
TATTAATCGATTGGTCTTTCGTTTCTCAGTCAATAGGTGGTCTCTTTAATGTGATTATGGCAGGTGGTCCCATGCTTTTGTGTTT  
TTTGATGAAAAAGTTATTACGATTGGCAATGATTCCAGGCTTATTTATGATGTTCCGGGGCATGATTTTACGATTAGAGACGA  
GGTCAAGGCTTTAAGGCTAAATGAGACCTATAGACCTTTGATTAG

## 4158.3 (SEQ. ID. NO. 274)

ATGTCCTCTCTTTCCGGATCAAGAATTAGTAGCTAAAAACAGTAGAGTTTCGTCAGCGTCTTTCCGAGGGAGAAAGTCTAGACG  
ATATTTTGGTTGAAGCTTTTGTGTGGTGGTGAAGCAGATAAGCGGATTTTAGGGATGTTTCTTATGATGTTCAAGTCAATG  
GGAGTTATGTTCAATGCTATGGAAATGTTGCTGAGATGAATACGGGGGAAGGTAAGACCTTGACAGCTACCATGCTGTCT  
ATTTGAACGCTTTTTCAGGAGAAGGAGTGATGGTTGTGACTCCTAATGAGTATTTATCAAAGCGTGATGCCGAGGAAATGGG  
TCAAGTTTATCGTTTTCTAGGATTGACCATTTGGTGTACCATTTACGGAAGATCCAAAGAAGGAGATGAAAGCTGAAGAAAAAG  
AAGCTTATCTATGCTTCGGATATCATCTACACAACCAATAGTAATTTAGGTTTGGATTATCTAAATGATAACCTAGCCTCGAA  
TGAAGAAGGTAAGTTTTTACGACCGTTTAACTATGTGATTATTTAGTGAATTTGATGATATCTTGCTTATAGTGCACAACTC  
CTCTGATTATTGCGGGTCTCCTCGTGTTCAGTCTAATTACTATGCGATCATTGATACACTTGTAAACACCTTGGTTCGAAGGA  
GAGGATTATATCTTTAAAGAGGAGAAAGAGGAGGTTTGGCTCACTACTAAGGGGGCCAAAGTCTGCTGAGAAATTTCTTAGGGA  
TTGATAAATTATCAAGAAGGAGAGCATGCGTCTTTTGTCTGCTACTGTTTATGCGATTTCGAGCTCAATAAGCTTTACTAA  
GATAAGGACTATATCATTTCTGGAATGAGATGGTACTGGTTGATAAGGGAACAGGGCGTCTAATGGAAATGACTAACTTC  
AAGGAGGTCTCCATCAGGCTATTGAAGCCAAGGAACATGTCAAATTTATCTCTGAGACGCGGGCTATGGCCTCGATCACCTA  
TCAGAGTCTTTTAAAGATGTTTAAATAGATATCTGGTATGACAGGGACAGGTAAGGTGCGGGAAGAAAGAGTTTATTGAACT  
TACAATATGCTGTAGTACGATTCACCAATCGTCCGAGACAACGGA  
TTGACTATCCAGATAATCTATATACCTTTACCTGAAAAAGTGTATGCATCCTTGGAGTACATCAAGCAATACCATGCTAAG  
GGAAATCCTTTACTCGTTTTGTAGGCTCAGTTGAAATGTCTCAACTCTATTCGTTCTCTTGTTCGTGAAGGGATTGCCCAT  
AATGTCCTAAATGCTAATAATGCGGCGCGTGAGGCTCAGATTATCTCCGAGTCAGGTACAGATGGGGCTGTGACAGTGGCTA  
CCTCTATGGCAGGACGTGGTACGGATATCAAGCTTGGTAAAGGAGTGCAGAGCTTGGGGGCTTGATTGTTATTGGGACTGA  
GCGGATGGAAGTACGCGGATCGACCTACAAATTCGTGGCCGTTCTGGTCTGTCAGGGAGATCCTGGTATGAGTAAATTTTT  
GTATCCTTAGAGGATGATGTTATCAAGAAATTTGGTCCATCTTGGGTGCATAAAAGTACAAAGACTATCAGGTTCAAGATA  
TGACTCAACCGGAAGTATTGAAAGGTGCTAAATACCGGAACTAGTCGAAAAAGGCTCAGCATGCCAGTGATAGTGTGGAC  
GTTACGACGTCGTCAGACTCTGGAGTATGCTGAAAGTATGAATATACAACGGGATATAGTCTATAAAGAGAGAAATCGTCT  
AATAGATGGTTCTCGTACTTAGAGGATGTTGTTGTGGATATCATTTGAGAGATATACAGAAGAGGTAGCGGCTGATCACTAT  
GCTAGTCGTGAATTATTTGTTTCACTTTATGTTGACCAATATTAGTTTTCATGTTAAAGAGGTTCCAGATTATAGATGTAAC  
GACAAAACCTGCAGTTCGTAGCTTTATGAAGCAGGTGATTGATAAAGAACTTTCTGAAAAAGAAAGAAATTACTTAATCAACATG  
ACTTATATGAACAGTTTTTACGACTTTCACTGCTTAAAGCCATTGATGACAACCTGGGTAGAGCAGGTAGACTATCTACAACAG  
CTATCCATGGCTATCGGTGGTCAATCTGCTAGTCAGAAAAATCCAATCGTAGAGTACTATCAAGAAGCCTACGCGGGCTTTG  
AAGCTATGAAAGAACAGATTATGCGGATATGGTGCATATCTCTGATGGGGCTGGTTGAGGTCACTCCAAAAGGTGAAAT  
CGTGACTCATTTTCCATAA

## 4158.4 (SEQ. ID. NO. 275)

ATGATAGGGACTTTTCGCGCTGCTCTTGTAGCTGTACTAGCAAATTTTCATCGTCCCTATTGAAATTACCCCAAAATAGTGCCAA  
TACTGAAATTCACCACCAGATGGGATTGGGCAGGTTCTCAGCAACCTCTTGCTCAAACCTGGTTGACAACCCAGTCAACGCC  
CTGCTTACTGCTAACTATATTAGAATCTTATCTTGGGCAGTCATTTTGAATCGCTATGAGAGAAGCCAGTAAAAATAGTCA



AGAAATTGCTAAAACTATCGCTGACGTGACTTCTAAAAATTGTCGAATGGATCATCAATCTGGCTCCATTGGAATCCTTGGCT  
TTGTTTTTAAAAACCAATTTCTGACAAGGGAGTCGGAAGCCTTGCCAACCTACGGTATTTTATTGGTTCTATTAGTAACGACTATG  
CTTTTTTGTGGCCCTGTGGTCAACCCCTTTGATTGCCTTCTTCTTTATGAGACGCAATCCTTACCCTCTAGTTTGGAACTGCCTCC  
GTGTCAGCGGTGTGACAGCCTTTTTCACTCGTAGTTCTGCGACTAACATTCCTGTCAACATGAAACTCTGCCATGACCTTGGGA  
CTCAACCCAGATACCTATTCTGTTTCTATCCCACTCGGTTCTACTATCAATATGGCTGGAGTAGCGATTACCATTAACCTTTTG  
ACCCCTGCTGCAGTTAACACTCTTGGAAATCCTGTTGACTTTGCCACAGCCTTTGTCCTCAGTGTGGTAGCAGCTATCTCATCC  
TGTGATGCTTCAGGTATTGCCGGAGGTTCCCTCCTTCTTATCCCACTGCTTGTAGCCTTTTCGGTATTTCTAACGATATTGCC  
ATACAAATTGTGGGTGGTTTGTGATTGGTGTCTATCCAAGACTCATGTGAAACAGCCCTTAACCTCTTCTACAGATGTCCT  
CTTTACC GCCGTGCGGAATACGCAGCAACCCGTAAAAAATAA

## 4158.5 (SEQ. ID. NO. 276)

ATGTCTATTAGCCAACGTACGACCAAGCTCATCTTAGCTACCTGTCTTGCTGCCTGCCTGCTTGTATTCTCAATCTTTCGTCA  
GCAGTTTCGGCTGGAATTATCGCTCTCTTGAGCCTATCTGATACGCGTAGAAGTACTTTAAAACTGGCTCGCAATCGTCTTTTT  
TCTATGCTTCTAGCTCTGGCTATCGGTGTTCTAGCTTTTCACTTGAGCGGATTTCATATCTGGAGTCTCGGCCTCTATCTGGCC  
TTCTACGTTCTTTAGCCTACAAGATGGGCTGGGAAATGGCATTACACCAAGCACTGTTTTGGTTAGCCATCTCTTGGTTCA  
AGAGACTACTAGCCTTCCAGACCTTCTAGTCAAGTAACTTCTTCTTGTGATTGGTTACAGGATTGGCTTGTGTTAACTCTTA  
TATGCCCTTACAGAGAAGAGGAAATCCAGCACTACCACACGCTGGTGGAAAGAAAAGTTAAAAGATATCCTCCAGCGCTTCAA  
ATACTATTTATCCAGAGGAGACGGACGCAACCGAGCACAGCTGGTAGCAGAATTAGACACGCTTTTGAAGAAGCCCTCAG  
ACTGGTCTATTTGGATCACTCTGACCACCTTTTACCAGACAGACTACCATATCCACTACTTTGAGATGAGACAGCGCAAAA  
GTCGTATCCTGAGAAAATATGGCCCAACAGATTAACACTTGTACCTTGCCGCCAGTGAAAGCCTGATCTTAGCGCAACTCTTT  
TCAAAAATTGCAAGGTCAACTGAGCCAGACCAATCCTGCTTCTGATTGCTAGATGAAATTGAACGTTATCTGGAAGTCTTCCG  
GAACCGCAGTCTGCCAAGACAAGAGAAGAAATTTGAAACCCGCGCCACCCTTCTTCAACTCCTACGTGAAGCCAAAACCTTC  
ATCCAAGTAAAAGTTGATTTTTACCAAAAATATAGACAGTAA

## 4158.6 (SEQ. ID. NO. 277)

ATGGAATCATGTGCTGCGATTGCTGTTTTTGGCGTCATCATTTGGTTTTAGTCATTGGATATGTCAGCATCTCAGCTAAGATG  
AAATCATCTCAGGAAGCTGCAGAGTTGATGCTTTTAAATGCTGAACAAGAAGCAACTAATTTACGTGGACAAGCTGAGCGTG  
AAGCGGATTTACTTGTAAATGAAGCCAAACGTGAAAGCAAGTCTCTTAAAAAAGAAGCACTATTGGAGGCCAAAGAAGAAG  
CCAGAAAATACCGTGAAGAAGTGGACGCTGAATTCAAAATCAGAACGTCAAGAACTCAAAACAAATCGAAAGTCGTTTGACAG  
AGAGAGCTACTAGCCTTGACCGTAAGGACGACAATTTGACGAGTAAAGAACAAACACTTGAACAAAAAGAACAAGTATTT  
CTGATAGAGCGAAAAACCTTGATGCGCGTGAAGAGCAATTAGAGGAAGTGAAGACAAAAAGAAGCAGAACTAGAGCGT  
ATTGGTGGCGTGTCTCAGGCAGAAAGCAGAGATATTATCTTGGCTCAGACAGAGGAAAACCTTGACCAGGGAGATTGCCAGTC  
GCATTGCGGAAGCTGAGCAAGAGGTCAAGGAACGTTCTGACAAAAATGGCCAAGGACATCTTGGTTCAAGCTATGCAACGTAT  
CGCTGGTGAATATGTAGCGGAGTCAACAACTCAACAGTTCTCTGCCAGACGATACTATGAAGGGACGCAATTATGGTCTG  
GAAGGTCGTAACATTCGTACCTTTGAAAGTTTGACAGGGGTGATGTGATTATCGACGATACACCAGAAGTGGTGACCTTGT  
CAGGATTTGATCCGATTGCTGCTGAGATTGCCCGTATGACTATGGAATGTTGCTCAAAGATGGTCTATACATCCAGCTCGT  
ATCGAAGAGTTGGTTGAGAAAAACCGTCAAGAGATTGACAATAAGATTCTGTAATACGGTGAGGCTGCTGCCTATGAAATTG  
GTGCGCCAAACCTTCATCCAGACTTGATGAAGATTATGGGACGTTTGCAGTTCCGTACTTCATATGGACAAAAATGTTTTGCGC  
CATTGATTTGAGGTTGCTAAGTTGGCTGGTATCATGGCGAGCGAAGCTTGGTGAATAAGCGCTCTTCCCGCTGCTGCTGGATT  
CCTTCACGATATCGGGAAGCCATTGACCATGAGGTGAAGGTAGCCACGTTGAAATCGGTATGGAATTGGCCCGTAAGTAC  
AAGGAACCCCAAGTTGTGGTGAATACGATTGTGATTGACACCGAGAGATTGGAAGCTGAGAGCGTGATAGCGTTATGCTGCG  
CTGCAGCAGATGCCTTGAGCGCAGCCCGTCCAGGTGCTCGTAGTGAGTCTCTTGAAGCTACATCAAGCGTCTCCATGATTG  
GAAGAAATTGCTAACCGCTTTGAAGGAGTGCAAACTAGCTTTGCCCTTCAAGCAGGACGTTGAAATTCGTATCATGGTCAATC  
CAGGAAAAATCAAGGACGACAAAGTCAAACTTGGCTCACAAGTTCTGTAAGAAAATTGAAAACAATCTCGATTATCCAG  
GAAATATCAAGGTAACCGTGATTGCGGAGCTTCTGTCAGTAGATTATGCTAAATAA

## 4158.7 (SEQ. ID. NO. 278)

ATGATGTTAAAAACCTCTATTGATACCTTGCTCGACAAGGTTCTTCAAAAATATTCACCTCGTAATCTTGGAAAGCAAAAACGTGC  
CCACGAATTGGAAGCAGGTGCCCCAGCAACTCAAGGTTTCAAGTCTGAAAAATCAACTCTTCGCGCTTTAGAAGAAATCGAA  
TCAGGAAACGTTACAATTCACCCAGATCCAGAAGGAAAAACGTGAAGCAGTGCCTGCGCGTATCGAAGAAAGAAAAACGCCG  
AAAGAAGAAGAAGAAAAAGAAAATCAAAGAGCAAAATTGCTAAAGAAAAAGAAGATGGTGAAAAAATTTAA

## 4161.1 (SEQ. ID. NO. 279)

ATGTCAGCATATCAATTACCGACCGTATGGCAGGATGAAGCTAGTAATCAAGGAGCTTTTACGGGGCTAAACAGACCAACAG  
CAGGTGCCCGTTTGAACAAAACTTGCCAAAAAGGAGAACAAGCTTTTCAAGCTTTTACTGGAACACCAAAATGGTGTGAA  
GGTACTATCTTATTGGAAGAAATTAAGAGCTGGTTTAAAGGAAGCGGCTTACGACTTGATATAAGATTGCTATCATGGATG  
GGGATCAATTCGGATCAGACTTTGTGAAGCTCAATCCAAATTCGAAGATTCCAGCCTTATTGGACCAAGTCAGGTACTGAAAA  
CGTAAGAGTCTTTGAGTCTGCTCATATTCTTTTACCTTGCTGAGAAAATTTGAGCCTTTTTACCAAGTAATCCTGTGGAAAA  
GGTAGAAGTTTGAATTGGCTATTCTGGCAAGCAGGTGCAGCACCTTTTCTAGGTGGGGGATTGGACATTTCTTCAATTATG  
CTCCTGAAAAATTTGAATATCCTATTAACCGTTTTACGATGGAAGTGAACGCCAGTTGGATTATTGGATAAGGAATTGGCT  
CAGAAACCTTATATTGCAGGCAATGACTATACGATTGCAGATATTGCTATCTGGTCTTGGTATGGACAGTTAGTTCAAGGAA  
ATCTTTACCAAGGTTCTGCAAAATTTCTGGATGCCTCAAGTTATCAAAAATCTAGTAAAATGGGCAGAAAAAATTGCCAATCG  
TCCAGCTGTTAAGCGTGGCTTGGAAGTAACCTTATACAGAAATTAATAG

## 4161.2 (SEQ. ID. NO. 280)

TTGGCAAGCTTGATCACTTCTATCATCATGTTCTATGTGGTTTTGATGTTCTAAGAGATACCAATTCAAAAAGATTCTCAGTCGG  
GAAGAACCGGTCATTGATCCTCTTGGTGCAACTCTAGGAATCAATTTCTGACGCGATTATGTTGTGGTCTATCTCTACAATAC  
TCGCTCAGTAAGAAATCCAATCCAATGCGCTGAAGGCAGCTGCTAAGGACAATCTTTCTGACGCTGTTACCTCACTTGGGA  
ACCGCAATGGCCATCCTAGCTAGTAGTTTCAATTAATCCGATTGTGGATAAACTGGTTGCTATCATCATCACTTTCTTTATCTTG  
AAGACTGCCTATGATATCTTCATCGAGTCTTCTTTAGTCTTTCAGATGGCTTTGACGACCGCCTGCTCGAGGACTACCAAAA  
GGCTATCATGGAATTTCCCAAAATCAGCAAGGTCAAAATCGCAAAGAGGTGCGACCTACGGTAGCAACATCTACCTGGATATT  
ACACTAGAGATGAATCCTGACTTGTCTGTTTTGAAAGCCATGAAATCGCGGATCAGGTGAGTCTATGCTGGAGGAGCGTT  
TTGGCGTCTTTGATACCGGATGTCCATATCGAACAGCACTTAACTGAGGATGAAATTTAGACAATGTCTGATAAAAATTG  
CTTATGCGTGAACAATTGATTGACCAAGGAAACCAACTAGAAGAACTCTTGACTGATGATTTGTCTATATTGCCAAGATG  
GAGAGCAGATGGATAAAGAGGCTTATAAGACCAAAAAAGAGTTAAATCTGCTATCAAGGACATTCAAATTACTTCCATCAG

TCAAAAAACCAAACCTCATCTGCTATGAGTTAGATGGTATCATCCATACCAGTATCTGGCGTCGCCACGAAACCTGGCAAAAT  
ATCTTTCATCAAGAAACCAAAAAAGAATAG

4162.1 (SEQ. ID. NO. 281)

ATGACAATTAACCTAGTACGAACGGATATGGACGGAACCTTCCTAGATGGGAATGGACGCTTTGATATGGATCGTCTCAAGT  
CTCTCTTGGTTTCTACAAGGAAAAAGGGATTTACTTTGCGGTAGCTTCGGGTCGGGGATTCTGTCTCTAGAAAAATTATTT  
GCTGGTGTTCGTGATGACATTATTTTCATCGCGGAAAAATGGCAGTTTGGTAGAGTATCAAGGTCAGGACTTGTATGAAGCGA  
CTATGTCTCGTGACTTTTATCTGGCAACTTTTGAAGGCTGAAAACTTCACCTTATGTAGATATCAATAAACTGCTCTTGACG  
GGTAAGAAGGGTTCATATGTTCTAGATACGGTTGATGAGACCTATTTGAAAGTGAGTCAGCACTATAATGAAAAATATCCAAA  
AAGTAGCGAGTTTGAAGATATCACAGATGACATTTTCAAATTTACAACCAACTTCACAGAAGAAACGCTGGAAGATGGGG  
AGGCTTGGGTAACGAAAAACGTTCCCTGGTGTAAAGGCCATGACAACTGGCTTTGAATCCATTGATATGTTCTGGACTATGTG  
GATAAGGGAGTGGCCATTGTTGAATTAGTTAAAAAACTTGGTATCACAATGGATCAGGTCATGGCTTTTGGAGACAATCTTA  
ATGACTTACATATGATGCAGGTTGTGGGACATCCTGTAGCTCCTGAAATGACAGCACTGAAATTTTAGAATTAGCAAAGAC  
TGTGATTGGTCACCATAAGGAACGGTCGGTTATAGCTTATATGGAGGGCTTATAA

4162.2 (SEQ. ID. NO. 282)

ATGGCAGATATAAAATTGATTGCATTGGACTTGGACGGGACCTTGCTGACTACTGATAAAAGGCTGACGGATCGTACCAAGG  
AAACCTTGCAAGCTGCGCGTGATCGTGGTATCAAGGTCGTATTGACAACCTGGTCGCTCCCTTAAAAGCCATGGATTTCTTCTC  
CATGAGTTAGGGACTGACGGTCAGGAAGATGAGTATACCATTACTTTAATGGTGGATTAGTTTCAGAAAAATACAGGAGAAAA  
TCCTTGATAAAACAGTCTTTTCATATGATGATGATGGCACGTTTGTATGAAGAAACAGAGAAATATCACTGCTCTTGTATGCC  
ATCTCAGAAGGAACAGTTTATCAAATCCAATCGGACCAAGAAAGTCTTTATGCCAAATTCATCCAGCTTTGACCTTTGTTC  
AGTGGACTTTGAAGACTTATCTAGTCAAATGACCTACAACAAATGCGTGACTGCTTTGCTCAAGAACCTTGGATGCAGCC  
ATTTCAGAAGATTCTCCAGAATTGTTTGACCAATATGAAATCTTTAAATCAGTGAAATGTTGCTAGAATGGTCACCAAAGA  
ATGTTTCAATAAGCAACAGGTTTGGCAAACTAATCAGCCATCTTGAATCGACCAAGTCAAGTGATGGCTTTGTGGTGACGA  
GGCCAATGACCTCTCTATGATTGAATGGGCAGGTCTTGGTGTGCTATGCAAAACGCTGTTCCCTGAAGTAAAGGCAGCCGCA  
AATGTAGTGACGCCGATGACCAACGATGAGGAAGCTGTGCGCTGGGCTATCGAAGAATATGTGCTAAAGGAGAACTAA

4164.2 (SEQ. ID. NO. 283)

ATGGAAAGTTTACTTATTCTATTATTAATTGCCAATCTAGCTGGTCTCTTTCTGATTTGGCAAAGGCAGGATAGGCAGGAGAA  
ACACTTAAGTAAGAGCTTGGAGGATCAGGCAGATCAATTTGTCAGACCAAGTTGGATTACCGCTTTGACCAAGCCAGACAAGCC  
AGCCAGTTAGACCAAAAAAGATTGGAAAGTGGTTGTCAAGCAGCGTTTGAAGAAGTGCGGATTGAATTGCACCAAGGCTGTA  
CCCAAGTCCGTCAGAAATGACAGATAATCTCCTCCAACTAGAGACAAGACAGACCAACGCTCTCCAAGCCTTGACGGAATC  
AAATGAGCAACGTTTGGAAACAAATGCGCCAGACGGTCGAGGAAAAAACTAGAAAAGACCTTGACAGACACGCTTACAGGCTTC  
CTTTGAGACAGTTTCTAAACAACCTGGAGTCTGTCAATCGTGGCCTTGGAGAAATGACAGACAGTTGCCCTGATGTGCGGAGCT  
CTTAACAAGGTTCTCTCTGGAACCAAGACGCGAGGGATTCTGGGAGAATTGCAACTGGGGCAAATATTGAAGACATCATGA  
CACCTGCCAGTACGAACGAGAATACGCAACGGTTGAAAACCTCTAGTGAACGAGTGGAGTATGCCATCAAGTTACCCGGAC  
AAGGGACCAAGAATACGTCTATCTGCCAATTGACTCTAAGTTTCCACTGGCAGATTATTACCGCTTGAAGAAGCCATATGA  
GACAGGTGACAAGGATGAGATTGAACGCTGTGCTAAGTCACTCCTAGCAAGCGTCAAGCGCTTTGCTAGGGATATTAGGAAC  
AAGTACATAGCACCACTCGGACGACCAATTTTGGAGTTTGTGTTGTTCCGACAGAAGGTCTCTACTCAGAAATCGTCCGCAA  
TCCGCTCTCTTTGATGATTTGAGACGGGAAGAACAGATTATTGTTGACAGGACCAAGTACCCTATCAGCCCTTCTTAACCTCCC  
TATCAGTTGGTTTCAAGACCCTTAATATCCAAAAGAGTGCCGACCATATCAGCAAGACTCTTGCCAGTGTCAAGACCGAGTTT  
GGCAAGTTTGGTGGTATTCTGGTCAAGGCACAAAAACATCTCCAACATGCCTCTGGCAATATTGATGAATTATTAACCGTCT  
GTACCATAGCTATCGAGCGGACGCTCCGTCACATTGAGTTGTGAGAAGGTGAGCCTGCGCTTGATCTACTCCATTTTCAAGAA  
AATGAGGAAGAATATGAAGATTAG

4164.3 (SEQ. ID. NO. 284)

ATGAAGATTAGTCACATGAAAAAAGATGAGTTATTTGAAGGCTTTTACCTAATCAAATCAGTGACCTGAGGCAAACCTCGAG  
CTGGGAAAAACTACCTAGCCTTACCTTCCAAGATGATAGTGGCGAGATTGATGGGAAGCTCTGGGATGCCCAACCTCATAA  
CATTGAGGCCCTTACCAGGATTAAGGTTGTCCACATGAAAGGACGCGGAGAAAGTTTATAACAATACCCCTCAAGTCAATCAA  
ATTACTCTCCGCTGCTCAAGCTGGTGAACCCATGACCCAGCTGATTTCAAGGTCAAGTCAACAGTTGATGTCAAGGAAA  
TTCGTGACTACATGTCGCAATGATTTTCAAATGAAAAATCCTGTCTGGCAACGGATTGTCCGAAATCTCTACACCAAGTAT  
GATAAGGAATCTACTCTCTACAGCTGCCAAGACCAACCACTGCTTTGAAACGGGCTTGGCCCTATCATACGGGACCA  
TGGTGGCTTTGGCAGACGCTATTAGCGAAGTTTATCCTCAGCTCAATAAGAGCCTGCTCTATGCGGGGATTATGTTGCATGAC  
TTAGCTAAGGTCATCGAGTTGACGGGGCCAGACCAGACAGAGTACACAGTGCGAGGTAATCTTCTTGGACATATCGCTCTCA  
TTGATAGCGAAATTAACCAAGACAGTTATGGAACCTCGGCATCGATGATACCAAGGAAGAAGTCGTTTTGCTTCGTCATGTCT  
CCTCAGTCAACACGGCTTGCTTGAAGTATGGAAGCCAGTCCGTCACGCATTATGGAAGCAGAGATTATCCATATGATTGAC  
AATCTGGATGCAAGCATGATGATGATGTAACAGCTCTTGTCTTGGTGGATAAAGGAGAGATGACCAATAAAATCTTCGCTA  
TGGATAATCGTTCTCTATAAACCAGATTTAGATTAA

4166.2 (SEQ. ID. NO. 285)

ATGAGTGAAAAAGCTAAAAAAGGGTTTAAGATGCCTTCATCTTACACCGTATTATTGATAATCATTGCTATTATGGCAGTGTCT  
AAGTTGGTTTATCCCTGCGGGGGCCTTTATAGAAGGATTTTACGAGACTCAGCCTCAAAATCCACAAGGGATTGGGATGTCC  
TCATGGCACCGATTGGGGCTATGCTAGGTACTCATCCAGAGGAAGGTTGCTCATTAAAGAAACGAGCGCAGCGATTGATGT  
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GAAGAAGTATAAGGGCCGCGAAAAAATGTTAATTTTGGTACTGATGCCTTTGTTTGGCCTCGGTGGTACAACCTTATGGTATGG  
GTGAAGAAACAATGGCCTTCTATCCACTCCTTGTGCCAGTTATGATGGCCGTTGGTTTGGATAGCTGACTGGTGTGCAATT  
ATTTTGTCTCGGTTCTCAAATCGGCTGTTTGGCATCTACTCTGAATCCATTTGCGACAGGTATTGCTTCAGCGACTGCGGGAGTT  
GGTACAGGGGACGGTATCGTACTTCTGCTGATCTTCTGGTTACCTTGACTGCTCTTACTGTTGTTTACCGTTATGCG  
GATAAGATTCAAAAAAGATCCGACTAAGTCACTGGTTTATAGTACTGCAAGAAGATTGAAACACTTAAACGTAGAAGAA  
CTTCTATCTGTAAGATTACACTTAGCAGCAAAACAAAAATCAGTTCTCTTCTTATTGTTGTTGACATTCTTGTATGGTATTGA  
GCTTCATTCCATGGACAGACCTTGGCGTTACCATTTTTGTATGACTTTAATACCTGGTTGACTGGTCTTCCAGTTATTGGTAATA  
TTGTCGGTTCTACTTCTGCACTAGGTACTTGGTACTTCCCAAGGGCGCAATGCTCTTGGCCTTATGGGTATCTCGTATTG  
GTGTTATTATGTTCTTAAAGAAAGATAAGATTATCTCTCTTCAATGAATGGTGCTGCTGACTTGGCTCACTGTTGCTTATGCTG  
TAGCGATTGCTCGTGGTATTCAAGTTATCATGAACGACGGTATGATTACCGATACAATCTCAACTGGGGTAAAGAAGGCTT



GAGCGGTCTATCTTCACAAGTCTTTATCGTTGTAACCTTATATCTTCTATCTACCTATGTCATTCTTGATCCCATCTTCATCTGGT  
CTTGCCAGCGCAACTATGGGTATCATGGC  
TCCACTTGGAGAAATTTGTAATGTCCGTCCTAGCTTGATTATCACTGCTTACCAATCTGCTTCAGGTGCTTTGAACTTGATTGC  
ACCAACATCTGGTATTGTGATGGGAGCTCTTGCACTTGGACGTATCAACATTGGTACTTGGTGGAATTCATGGGCAAACTCG  
TAGTCGCTATTATTGTAGTGACCATCGCCCTTCTTCTCCTTGGAACTTCCCTTCCATTCTATAA

4166.3 (SEQ. ID. NO. 286)

ATGAAAATAGATATAACAAATCAAGTTAAAGATGAATTTCTTATATCATTAAAAACCTTGATTTCTATCTTCAGTACTCAA  
TGAAGGAGAAAAATGGAACACCTTTTGGACAAGCAATCCAAGATGTCCTAGAAAAAACTTTAGAGATTTGTGCGAGACATAGG  
TTTCACTACCTATCTTGACCCTAAAGGTTATTACGGATATGCAGAAATCGGTGAGGAGCAGAGCTTCTGGCCATTCTCTGTC  
ATTTGGATGTTGTTCCATCAGGTGATGAAGCAGATTGGCAGACACCGCCATTGGAAGCAACTATCAAAGACGGCTGGGTATT  
CGGACGTGATGTTCCAAGATGATAAAGGCCCTTCGCTCGCAGCTCTGATGCAAGTAAAAAGCTTGCTGGACCAAGGTATTGAG  
TTCAAAAAGCGCGTACGCTTTTATCTTTGGTACCGATGAGGAAACCTCTGGCGCTGCATGGCAGCTACAATACCATCGAAG  
AACAGGCCAGTATGGGCTTTGACCTGACTCATCTTTTCTCTGACCTATGCTGAAAAAGGGCTTCTACAGGTCAAACCTTCAT  
GGCCCTGGATCGGATCAACTAGAGCTTGAAGTAGGAGGCGCCTTAACGTTGTACCAGACAAGGCCAACTACCAAGGTCTCC  
TCTATGAACAGGTTTGTAAACGGTCTCAAAGAAGCTGGTTATGATTACCAAACTGAAACAAACCGTAACGGTTCTCGGAGT  
GCCAAAGCATGCTAAGGATGCTAGTCAAGGTATCAATGCTGCTACCGACTAGTACCATTCTTGCTCCTCTCCAAGAACACCG  
CTGCTCTCAGTTTTCTTGCAACACAAGCAGGTCAAGACGGCACAGGAAGACAAATCTTTGGTGATATAGCAGATGAACCTTC  
TGGTCACTATCTTTAATGTGCGAGGTCTCATGATCAATCATGAACGTTCTGAAATCCGTATTGACATTCCGACTCCTGTCTT  
AGCTGACAAGGAAGAACTAGTAGAGTTGCTTACAAGATGTGCACAAAACCTACCAACTCCGCTACGAAGAGTTTGACTATCTA  
CGCCTCTATACGTCGCAAGACAGTAAACTGTTAGCACACTGCAAAATCTACCAAGAAAAAGCTGGCGACTGCGAAGACAGC  
CTGCTATTTTATCCGGTGGTGCCACTTTTCTGCTCGCACCATGCCAAATTGTGTAGCCTTCGGCGCCTTATTCCAGGAGCGAAG  
CAGACAGAACATCAGGCAAAATGAATGTGCCGTCTAGAAGATTGTACCGTGCTATGGATATTTATGCCGAAGCCGTCTATC  
GACTTGAACCTTAA

4169.1 (SEQ. ID. NO. 287)

ATGCTAATTCATTTGTCAAGTTGTTAGTCTCTCAATTTTGTCAAATTTAGCAGATATTTTCTTTAGAGTAACAATCATTGCT  
AACATACATATATTTCAAAAATCAGTAATTGCCACATCACTAGTTCCTATCTTAATAGGAATATCCTCTTTTGTGCGAGTCTT  
TTAGTTCCGTTGGTTACTAAAAGGTTAGCGCTAAAATAGGGTTTTATCTTTATCTCAATTTGGAAAGACTATATTATTGGCGAT  
ACTGGTAGGAAATGTTTACCGTAATGCAATCCGTAGCGCCTTTGGTGACCTATCTATTTGTTGTTGCAATTTCCATAGTAGT  
GTTTTGACGACCCGTTTCTATGCTATTGTGCCAGCTATGCGACCGATTGGGTAAGGCTAATTCAGCCTTATCAATGACT  
GGTGAAGCTGTTCAATTGATAGGTTGGGGATTAGGTGGACTCTTGTGTTGCAACAATTTGGTCTGTTACCTACCAGGTGATCAA  
TTTAGTCTGTATATCAATTTCTAGCTTTCTGATGTTATTTCTCTCAACGCTGAAGTGGAGGTGTTAGAGTCAGAACTAATCT  
TGAAATTTTGTCAAAAGGTTGGAAGTTAGTTGCTAGAAATCCTAGATTAAGACTTTTTGTATCAGCAAAATTTATTGGAAATTT  
TTTCAAAATCAGATTGGGTTTCTTCCATTATACTTGTGTTTGTAAACGGAGTTATTAAATAAAACGGAAAGTTACTGGGGATATT  
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AACCCTAATTTATCACCCTAAATCTAAAAACCATCCAGAATCCTTGCTTAGCTTAGATCCCTGGATGGTTTCTTTTTTCAACCCA  
ATGGGTGTTTTTACTAGACAAAAAAGAGTTTCCCTTTATGGTATAAGTGTAGAAAAAACACAAAAAGAAAGGAACTCA  
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CGCCGCTACTGTGCTTATTCGGATTAGATATCCTTGTCAGTTCTCTTTCAACTGTTAACAGGTTATGGAACGGACTATGCT  
TGTAAGAATTGTCAAGTGTGCTGATGCTACTT  
TCCAAAATTTGTTGGAAGGAGGGCAGCTTGCTTCACAGCCAACCTTATCCCGTTTTCTTTCCAGAACTGACGAGGAAACAGTCC  
ATAGTTTGGCGATGCCTCAACCTTGAATTGGTGAATTTCTTTTACAGTTTACCAAGCTAAACCAACTCATTGTAGATATCGATT  
CTACCCATTTTCAACATTTATGGCAAGCAAGAAGGTGTGCTTATAACGCCCACTATCGTGCTCATGGCTATCATCTCTTTAT  
GCTTTTCGAGGGGAAGACAGGTTATTGTTTCAATGCCAGCTTCTGCTCGTAAATCGTTATTGTTCTGAAGAGGCGACAGACTT  
TATCACACCTGTTTTAGAACGGTTAATCAACTTCTCTTTCGAATGGATAGTGGCTTTGCGACCCCAAAATATACGATTTAAT  
TGAAAAACAGGGCAATACTACCTCATAAACTCAAGAAAAAATCACTGTTCTGAGCCGCTCTGGAGACCTTTCCCTCCCTTGC  
CCACAGGATGAGGACTTAACCATCTTGCCCACTCGCCTACTCAGAAACTCTATCAAGCAGGACTTGGTGCACAAAGC  
GTCGTGTCTGCCAGTTCTCTGAACGAAAAGAAGGAACTTGTCTACGATGTTATTTCTCTCGTTACAAATATGACGAGTGGG  
ACAAGCCAAGACAGTTTACGCTTATCGTGGACGTGGTCAAGCCGAGAATTTATCAAGGAGATGAAGGAGGGGATTTTTTG  
CGGATAAAACGGATAGTTCAACCTTAATCAAAAAACGAAGTTCGTATGATGATGAGCTGTATCGCCTACAATCTCTATCTTTT  
CTCAAACTCTAGATGAGGTGACTTCCAAACTTTAACAATCAAAACGCTTCCGCCATCTTTCTTTCAGCTGGTGGGAAAAATG  
TGTTGCAACAGGACGCAAGCAGCTCCTCAAATTGTCTAGTCTCTATGCCTATCCGAATTGTTTTCAGCACTTTATTCTAGGAT  
TAGAAAAGTCAACCTGAATCTTCTGTTCTTATGAACCACCTAGAAGAAAAGCGTCGTTAATGATGCATTAA

4169.3 (SEQ. ID. NO. 288)

ATGATGGAGTTTTTCAACAGCTTCCTCATTTAGAGCCATATGGCAATCCTCAGTATTTTGTGTTATGTGATTGCTGCAACCTTG  
CCCATCTTTATAGGTCTCTTTTTCAAGAAACGCTTTGCCTGGTATGAAGTGTGGTAAGTCTCTCTTTATTGTACCAATGTTG  
GTGGGTGGAAGACCAATCAACTAGCTGCCTTGGGTATTACCTTTGTGCGGAAATATTGCTCCTGCTTTTCTCAAGCATTAA  
TCGAAAAAGCAAGGATGGCAAGTGGGTCTTCTACTTAGTTAGTTTCTGTCCCTACTCCGATTATCTTTGTCAAGGTGCAAC  
CAGCTATCAATGGAACGCAAGTCTTTGCTTGGGTTCTTGGGAATTTCTTACCTGACCTTTCGTTCCGTTGGAATTGTATCGAGC  
TGAGAGATGGAGTGATTAAAGGATTTACCTCTGGGAATTCCTCTCTCTTCATGCCAAGCTTTCTGAGTGGTGGTCCAA  
TCGATCGCTTTAAGCGATTTAATGAAAAATTACAGGCTATTCTGAGCGAGATGAGTTGATGGATATGCTGGATGAATCTGTC  
CGCTATATCATGTGGGGCTTTTTGTATAAGTTTATCCTAGCTCATGTTTATAGGAGAGACCTTACTACCTCCTCTGAAGAAATTA  
GCCTTGCACTCAGGTGGCTTCTTAACTCTATGCCTTGGCAGTTATGTATACTTTTGGTCTGGAACCTCTCTTTGACTTTGCA  
GGTTATTCTATGTTTGTCTTGGCCATCTCAAACTTTAACAATCAAAACGCTTCCGCCATCTTTCTTTCAGCTGGTGGGAAAAATG  
GATTTAAAGGAGTTTGGAAATCGCTGGCATATGAGTCTGTCTTCTGGTTCCGTGACTTTGTCTTTATGCGAATGGTGATGGT  
GTTAACCAGAAAGAAAGCTTTTAAAAATCGTAATGTAACCTCAAGCATGGCCTACATTGTAATATGCTGATTATGGGATTTT  
GGCATGGTGTGACCTGGTACTATATCGCCTATGGACTCTTCAAGGACTAGGCTTGGTCAATCAATGATGCTGGTGGTTCGCAAG  
AAAAAACGCTCAATAAGGAACGGAAGCAAGGGAAGGCTGCCCTACCTGAGAATCGCTGGATTAGTTGCTTGGCATG  
GTTGCTACTTTCCATGTTGTCATGTTGTCATTCTTAATCTTTTCTGGATTCTTGAATAATCTATGGTTTAAAAAATAA

4169.4 (SEQ. ID. NO. 289)

ATGCTTAAACGCTTATGGATGATCTTCGGACCGGCTTGATCGCTGGTTTGTGGTITTTCTGCTCATTTTCTTTATCCTACTG  
AGATGCATCATAACTAGGAGCTGAAAGCGTTACCGAGTGGCTACTACTATCGATAGTTTAAAGGACGGAGTCAAAAAAG  
CAGAGCACTATCGATGCAAAATGTGCGTTTGTCCCTCTTTGGCTCTAGTGAATGGCTCGTTTGGACGGTGCTCATCTGCG  
GGTATTAGCTGAGAAATACAATCGTTCCTACCGTCCTTATCTTTTAGGACAGGGGGGAGCTGCATCGCTTAACCAATATTTTG  
GAATGCAACAGATGTTTACCACAGCTGGAGAAATAAACAAGTTGTGATGTTATCTCACTCAGTGGTTCAGATAAAAAATGGCTA  
TGATCCAGACGCTTCCAGCAGTATTTTAAATGGAGACCAAGTTGACTAGTTTCTGAAACCTCAATCTGGGATCAGGCTAGTC  
AATATGCAGCGACTCGCTTACTGCAACAGTTCCTAAACGTAAGCTATGAAGGACCTGGTTCAGAAGTTGGCAAGTAAAGAAGA  
ATTGTCGACAGCAGACAATGAATGATTGAATTATGGCTCGTTTAAATGAACCCGAAGCTTCCTTTTGGTCAGTTTTCGG  
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AGAAGGATTTGAAGAAATTAAGGATTCTCAGAAAAGCTTTACCTATCTCAAGTCCGAGAGGATAATGACTTGCAGTTGGT  
TTTAAACACAGTTTCTAAATCTAAGGTAACCCGATTTTATCTATCCCACTGTTAATAAAAAATGGATGAACTATGCTGGTC  
TAGCAGAGGATATGTACCAACAACCGGTGCAGAAAGTTTCGCTACCAGTTAGAAAAGTCAAGGTTTTACCAATAATAGCAGATTT  
TTCTAAGGACGCGGGGAGCCTTTCTTTATGAAGGACACCATTACCTTGGTTGGTTGGGTTGGTTGGCTTTTGACAAGGCAG  
TTGATCTTTTCTATCCAATCCCACACCAAGCTCCGACTTACCATCTGAATGAGCGCTTTTTCAGCAAAGATTGGGCGACTTAT  
GATGGAGATGTCAAGAATTTCAATAG

4169.6 (SEQ. ID. NO. 290)

ATGGAGAAAAACCTCAAGGCTTTGAAACAAACAACAGACCAAGAAGGCCAGCAATTGAACCTGAAAAGGCAGAGGATAC  
CAAGACAGTCCTCAAAAATGTTTACTTCGAGGATGCAAGCTGTCAAGGACCGCACCTTGAGTGACTATGCAGGTAACCTGGCAATCA  
GTTTATCCTTTCCCTTGAAGACGGCCAGCTTTGACCAAGTCTTTGACTACAAGGCTAAGTTGACTGGTAAGATGACCCAGGCTGA  
GTACAAGGCTTACTATACAAAAAGGCTATCATACAGATGTGACTAAGATTAACATTACTGATAATACTATGGAATTTGTTCAA  
GGTGGACAAAGCAAGAAATACACTTACAAGTATGTCCGTAAGAAAAATTTGACTTACAAGAAAGGCAATCGTGGCGTGCCT  
TTCCTCTTTGAAGCCACAGATGCTGACGCTGGACAATTCAGTATGTTCAAGTTAGTGACCACAATGTTGCCCCAGTTAAGGC  
AGAACATTTCCCATATCTTCTTTGGAGGCCACAAGCCAAGGCCCTCTTTGAAGAAATGGACAACCTGGCCAACCTACTACCCA  
GATAACCTTACTGGCCAAGAAATCGCCCAAGAAATGTTGGCGCATTTGA

4170.3 (SEQ. ID. NO. 291)

ATGAAAGATGGTCATTGCTAGCCCATCATATTCGTTTGTTGAATGGGCGGATTTTTCAAAAGTTACTGAGTCAAGATCCTGA  
GGCTCTTTTATAGGGGTGAACAGGGCAAGATTGACGGCTTTTATGGAATAGTGAACCTGGCTGCGCAACTGCACAGATATC  
CGCTTTCGCACTGGACTTGGCAATAATACGCTTAGCGACTATGATAAAAAAGCTAGAGGAACAAAAGCTTGTAATTGTTAGTC  
CGTGTGGAAAAGACAAGCGTAAGAAGTATTTAGTTTTAACGGAGTTAGGCAAGTCCCAGAAAGAGTGGGGCATCGTGTC  
GTCAGAAATTGGATACTACTCTTTTACAAAGGATTTTCAGAGGAAGAAATTCACCAATTTGAAGGTTTTCAAGAAAGAATTTT  
GGCGAAATCTGAAAGAGAAGGGAATGAGGTTTAG

4170.4 (SEQ. ID. NO. 292)

ATGACTAATTTAATTGCAACTTTTCAGGATCGTTTTAGTGATTGGTTGACAGCTCTATCTCAACATTTGCAAGTGTGCGCTTTTG  
ACCTTGTTACTAGCTATTTTGCTTGCGATTCCCTTGGCTGTTTTCTTCGCTATCATGAGAAGCTGGCCGACTGGGTCTTGCGAG  
ATTGCAAGGTATTTTCCAGACCATCCCGTCTTGGCCTGTGTGGGGCTCTTTACCCCTTTGATGGGAATTTGGGACCTTCCGGCGCT  
TTGACAGCTCTAGTGATTATGCGATTTCCTCTATTTTGC AAAA TACTATCACTGGGCTGAAGGGAATTGATCCGAACCTGCA  
AGAGGCTGGGATTGCCTTTGGGATGACCAGATGGGAACGTCTCAAGAAATTTGAAATTTCCACTCGCCATGCCTGTTATCATG  
TCTGGGATTCGGAGCGGCAGCTGTTTTGATTATCGGTACGGCAACCTTGGCGGCGCTTGATTGGTGCAGGGGGAGTACGGTTCCTT  
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CTTATAGTCCAGCTCTTTTGGTTCAAAAAGAGAAGGAAAACTTGGTTATTGCTGGGAAAAATAGGTCAGAACCCAGAAATTTT  
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CTTTATGAAGCTCTGAAAAAAGCGCATATTGACATCTATCTGAAATTTACTGGTACGGTGACTGAAAGTTTGCTTCAACCATC  
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CTTATCAAGGGGCTCCACTCATGAAAGAAGCTCTTCTCAAGAAACACCCAGAGTTGGAAAGAGTCTTAATACATTGGCTGG  
TAAGATTACAGAAAGCCAGATGAGCCAGCTCAACTACCAAGTCGGTGTGGAAGGCAAGTCAGCAAAGCAAGTAGCCAAGGA  
GTTTCTCCAAGAACAGGTTTGTGTAAGAAATGA

4170.5 (SEO. ID. NO. 293)

ATGATGTCATACTTATTTGCAAAAAGAAAATTGAAAAATATCAAAACAACCCTAGGTGAAATGTCAGGTGGTTACCGTCGTATGG  
TTCGGGCTATGGCTGTTTATAGGATTTTCAGGAACCTATGAAGGCTATCTGGGATGACCTCTTTGCCCATCGTAGTTTTCGCCAG  
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GTTATTTACCTTATTTTGGCCCTACAGAAAGGCTTTTATGGTGAGGTGCTGACGACACTTTTATCTTCAACAGTCATGACGCCAA  
TGGACTTCTAGTTTGGATTTATCAGGCAAGTTTAAAGAAAGGAAAGGACGAGGATTTGTCGCGCTAAACTGGACGGCAAGGGC  
TGACAAAGATCTTTCCATTAGTGTGCTTTGGTGGTTGGCTTTGGCTTCATTTATCAGTCTATTGGTGCCAATCGTCCCTAT  
CGTGATTCAATCACAGATGCAACCAATGGGGTAGGGCAAATCCTCATGACAGCTGTTTACCGTGAAACAGTGGATATCTCTGGG  
CGGCTACCAATGTCTTTTCAATCTATCTCTGGTGGGGAGAAAGCCTGCAAAATCAAGGGAATAATCTAATTTATCTCATTAAC  
AGCTAGATTGGTTGGTATCAATGGAGCAAGGCAAGCTAAGCAGAATCTGATTACTTAAGT

4170.6 (SEQ. ID. NO. 294)

ATGAGAAATATGAAGGCCAAAAATATGCTGTTTGGGTGGCTTTTTCTTAAATTTGACTTATGCCATTGTTGAGTTTATTGCAGG  
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AGAAACAATCTCCAATCGTGAAGAAGACAATCAGTACACCTTGGGCTATAAGCGGTTTAGCTCTGCTAGGACCGCTTGGTAACA  
GCTGTGATTCTCGTAACGGGCTCTGTTCTAGTCATTTTGGAAAATGTCACGAAGATTTTGCATCCGCAACCAAGTCAATGATGA  
GGGGATTCTCTGGTTAGGAATTTATTGCGATTTACTATCAATCTGTTAGCGAGATCTGGTGGTTGGTAAGGGAAGAACAAGAAT  
GAGTCTATTCTGAGTCTGCATTTTCTGGAAGATACGCTAAGGGTGGGTAGCTGTTATCTCTGATGGCGATTTGTTCTTCGATTTC

GGACTGGTATATCCTAGATCCTCTTTTGTCCCTTGTCATTTCTTTCTTTATTTCTTTTCAAAAGCCCTTCCACGTTTTTGGTCTACA  
CTCAAGATTTTCTTGGATGCTGTGCCAGAAGGTCTTGATATCAAGCAAGTAAAGAGTGGCCTGGAGCGATTGGACAATGTGG  
CCAGCCTTAATCAGCTTAATCTCTGGACTATGGATGCTTTGGAAAAAATGCCATTGTCCATGTTTGTCTAAAAAGAAATGGAA  
CATATGGAACTTGTAAGAGTCTATTTCGAATTTTCTTAAAGATTGTGGTTTTCAAAATATTACCATTGAAATGATGCTGA  
CCTAGAACTACCAAACCCATAAGCGAAAGGTGTGTGACTTGGAAACGGAGTTATGAGCATCAACATTAG

## 4170.8 (SEQ. ID. NO. 295)

ATGATTGAATACAAAAATGTAGCACTGCGCTACACAGAAAAGGATGTCTTGAGAGATGTCAACTTACAGATTGAGGATGGG  
GAATTTATGGTTTTAGTAGGGCCTTCTGGGTCAAGTAAGACGACCATGCTCAAGATGATTAACCGTCTTTTGGAAACCACTGA  
TGGAAATATTTATATGGATGGGAAGCGCATCAAAAGACTATGATGAGCGTGAACCTTCGTCTTTCTACTGGTTATGTTTTACAGG  
CTATTGCTCTTTTCCAAATCTAACAGTTGCGGAAAAATATTGCTCTCATTCTGAAATGAAGGGGTGGAGCAAGGAAGAAATT  
ACGAAGAAAACAGAAGAGCTTTTGGCTAAGGTTGGTTTACCAGTAGCCGAGTATGGGCATCGCTTACCTAGTGAAATATCTG  
GTGGAGAACAGCAACGGGTGCGGTATTGTCCGAGCTATGATTGGTTCAGCCCAAGATTTTCTCATGGATGAACCCCTTTTCGGCC  
TTGGATGCTATTTTCGAGAAAACAGTTGCAGGTTCTGACAAAAGAATTGCATAAAGAGTTTGGGATGACAACGATTTTTGTAA  
CCCATGATACGGATGAAGCCTTGAAGTTGGCGGACCGTATTGCTGTCTTGCAGGATGGAGAAAATTCGCCAGGTAGCGAATCC  
CGAGACAATTTTAAAGCGCCTGCAACAGACTTTGTAGCAGACTTGTTTGGAGGTAAGTGTTCATGACTAA

## 4171.1 (SEQ. ID. NO. 296)

ATGTCAGCAGTTGCTATTTTCACTATGACCAAGGTTATGCAAGAAACCCACGGAAATCCTTCTAGTATTCATGGTCATGGTCTG  
TCAAGCTGGCAAACTCTTGGGAGAACCCGTCAGGAAGTACGCCAGTTACTAAGGACAAAACCTCAACATATCTTTTCACT  
TCTGGTGGGACTGAAGGCAATAATACTACCATCATTTGGTCTACTGTCTTCGTACCAAGAAACAGGAAAAACATATCATACAA  
CTGCCATCGAGCACCATGCTGTCTTGAACAATTGATTACTTGGTTCAACACTTTGGGTTTGAAGCAACCATTTATCCAGCCA  
GAAAATCAAGAAATCACAGCCAGCAAAATCAAAAGGCTTTACGTGACGATACGATTTTGGTTTCTACCATGTTTGTCAATA  
ATGAGACAGGAAACCTACTGCCATCGCTGAAATTTGGCCAAATCTCAAGCAACACCCTGCTCATCATGTTGATGCAAGT  
TCAGGCTATTGGTAAAATCCCAATTCATTGAGAAGAAATGGGCATTGATTTTCTCACTGCTTCTGCCACAAAATTCATGGTC  
CTAAGGGAATCGGTTTTCTCTACGCATCTAGCATTTGATTTTCTCTATCTACATGGCGGAGACCAGGAACAGAAAAAACG  
TGCAGGAACCTGAAAATCTGCCATGTCAGGCTGTTGTCAGGCTAAAAGAACCTAGAAAAACAAGAAAGCAATTT  
TCAACATGTACAAAATCTAGAAAATGCTTTCTGCGAGAGCTGGAGGGCACTTCAAGTATTACCTGAATGAGGAAAAACATCAT  
CTCCCTTATGTTCTCAATATTGGATTCTCTGGTCAGAAAAATGACCTCTTACTCCTTCGGCTAGATTTAGCTGGAATTTCAATC  
TCTACTGGCTCAGCCTGTACTGCAGGCGTTGTCCAATCCAGCCATGTTCTTGAAGCCATGTATGGCGCAAAATTCAGAACGCTT  
GAAGGAATCCCTTCGCATCAGTTTGTGCGCCAAAAATACCGTTGAAGACCTACAAACCCCTCGCAAAAACCTTAAAGAAAT  
ATCGGAGGTTAG

## 4172.1 (SEQ. ID. NO. 297)

ATGTTATTCAAATTATCTAAGGAAAAAATAGAGCTAGGCTTATCTCGTTTATCGCCAGCCCGTCGTATTTTTTTGAGTTTTGCC  
TTGGTCATTTTACTAGGCTCTCTTCTTTTGGAGCTTGCCCTTTTGCCAAGTTGAAAGCTCACGAGCGACTTATTTTGATCATCTTT  
TCATTGCTGTCTCTGCAGTCTGTGTGACGGGTCTCTCAACCCTTCCAGTAGCTCACACCTATAATATCTGGGGTCAAATAATC  
TGTTTGCTCTTGATTACAGATCGGTGGTCTAGGGCTCATGACCTTTATTGGGGTTTTCTATATCCAGAGCAAGCAAAAGCTTAG  
TCTTCGTAGCCGTGCAACTATTACAGGATAGTTTATGTTATGGAGAACTCGATCTTTGAGAAAAGTTTGCTATTCTATTTTCT  
CACGACCTTTTTGGTTGAGAGCTTGGGAGCTATTTTGCTTAGTTTTCGCCCTTATTCCTCAACTTGGCTGGGGACGTGGTCTTTT  
TAGTTCCATTTTTCTAGCGATCTCAGCCTTCTGTAATGCCGGTTTTGATAATTTAGGGAGCACCAGTTTATTTGCTTTTCAGAC  
CGATTTACTGGTCAATCTGGTGATTGCAAGCTTGATTATTACAGCGCGCCTTGGTTTTATGGTCTGGTTTGATTGGCTGGTCA  
TGTAAGGAAGAAAGAAAAAGGACGTCTGCACTTTTATACGAAGCTTGTAATTTTACTATAGGTTTGTGTTATTTGGAA  
CAGCAACTACTCTCTTTCTTGAGTGGAACAATGCTGGAACGATTGGCAATCTCCCTGTTGCCGATAAGGTTTTAGTTAGCTTT  
TTTCAAAACAGTGACGATGCGAACAGCTGGCTTTTCTACGATAGATTATACTCAGGCTCATCCTGTGACTCTTTTGATTTATATC  
TTACAGATGTTTTCTAGGTGGGGCACCTGGAGGAACAGCTGGGGGACTCAAGATTACGACATTTTTTGTCCTCTTGGCTTTTGC  
ACGAAGTGAGCTTCTAGGCTTGCCCTCATGCCAATGTTGCGAGACGAACGATCGCGCCGCGAACGGTTCAAAAATCCTTTAGT  
GTCTTTATTATCTTTTTGATGAGCTTCTTGATAGGATTGATTCTGTAGGGATAACAGCCAAAGGCAATCCTCCCTTTATCCAC  
CTCGTATTTGAAACCATTTACGCTCTTATGATCAGTTGGTGTAAACGGCAATCTGACTCTGACCTTGGGAAATTTGGCTCTCAG  
TGTTATCATGCCACTTATGTTTATGGG  
ACGAATTGGTCCCTTGACCTGTTTGTAGCTTGGCAGATTACCATCCAGAAAAGAAAGATATGATTCACTATATGAAAGCA  
GATATTAGTATTGGTTAA

## 4172.2 (SEQ. ID. NO. 298)

ATGTCAGATCGTACGAATTGGAATTTTGGGCTTGGGAATTTTTGGGAGCAGTGTCTAGCTGCCCTAGCCAAGCAGGATATGA  
ATATTATCGTATTGATGACCACGCAGAGCGCATCAATCAGTTTGAGCCAGTTTTGGCGCGTGGAGTGATTGGTGACATCAC  
AGATGAAGAATTATTGAGATCAGCAGGGATTGATACCTGCGATACCGTTGTAGTCGCGACAGGTGAAAAATCTGGAGTCGAGT  
GTGCTTGCCTTATGCACTGTAAGAGTTTGGGGGTACCGACTGTTATTGCTAAGGTCAAAAGTCAGACCGCTAAGAAAGTGC  
TAGAAAAGATTGGAGCTGACTCGGTTATCTCGCCAGAGTATGAAATGGGGCAGTCTCTAGCACAGACCATTTCTTTCCATAA  
TAGTGTGTAGTCTTTTTCAGTTGGATAAAAAATGTGTCTATCGTGGAGATGAAAATTCCTCAGTCTTGGCGAGGTCAAAAGTGA  
GTAAATTAGACCTCCGTGGCAATACAATCTGAATATTTGGGTTTCCGAGAGCAGGAAAAATCCCCATTGGATGTTGAATTT  
GGACCAGATGACCTCTTGAAAGCAGATACCTATATTTGGCAGTATCAACAACCAAGTATTTGGATACCCTAGTAGCATTTGA  
ATTCGTAA

## 4172.3 (SEQ. ID. NO. 299)

ATGAAGTTATTGTCTATCGCAATTTCTAGCTATAATGCAGCAGCCTATCTTCATTACTGTGTGGAGTCGCTAGTGATTGGTGG  
TGAGCAAGTTGGGATTTTGATTATCAATGACGGGTCTCAGGATCAGACTCAGGAAATCGCTGAGTGTTAGCTAGCAAGTAT  
CCTAATATCGTTAGAGCCATCTATCAGGAAAAATAAATGCCATGGCGGTGCGGTCAATCGTGGCTTGGTAGAGGCTTCTGGGC  
GCTATTTTAAAGTAGTTGACAGTGATGACTGGGTGGATCCTCGTGCCCTACTTGAAAAATCTTGAACCTTGCAGGAACTTGAG  
AGCAAAGGTCAAGAGTGATGTCTTTGTGACCAATTTGTCTATGAAAAGGAAGGGCAGTCTCGTAAGAAGAGTATGAGTT  
ACGATTCACTGTTCGCTGTCGGCAGATTTTGGCTGGGACCAAGTTCGGAATTTCTCCAAAGGCCAGTATACCATGATGCAC  
TCGCTGATTTATCGGACAGATTTGTTGCGTGCTAGCCAGTTCTAA

## 4172.4 (SEQ. ID. NO. 300)

ATGAAATTCAATCCAAATCAAAGATATACTCGTTGGTCTATTCCGCCGTCTCAGTGTCCGGTGTTCCTCAGTTGTTGTGGCTAG  
TGGCTTCTTTGTCCTAGTTGGTCAGCCAAAGTTCTGTACGTGCCGATGGGCTCAATCCAACCCCAAGGTCAAGTCTTACCTGAAG  
AGACATCGGGAACGAAAGAGGGTGACTTATCAGAAAAACAGGAGACACCGTTCTCACTCAAGCGAAACCTGAGGGCGTTA  
CTGGAATACGAATTCACCTCCGACACCTACAGAAAGAGTGAAGTGAGCGAGGAAACAAGCCCTTCTAGTCTGGATACAAAT  
TTTTGAAAAAGATGAAGAAGCTCAAAAAAATCCAGAGCTAACAGATGTCTTAAAGAAACTGTAGATACAGCTGATGTGGA  
TGGGACACAAGCAAGTCCAGCAGAACTACTCCTGAACAAGTAAAGGTGGAGTGAAAGAAAAATACAAAAGACAGCATCG  
ATGTTCTCTGCTGCTTATCTTGA AAAAGCTGAAGGGAAGGTCTTTCACTGCCGGTGTAAACCAAGTAATTCCTTATGAACATA  
TTCGCTGGTGATGGTATGTTAACTCGTCTATTACTAAAAGCTTCGGATAATGCTCCTTGGTCTGACAAATGGTACTGTAAAAA  
TCCTGCTTTACCTCCTCTTGAAGGATTAACAAAAAGGGAATACTTCTATGAAGTAGACTTAAATGGCAATACTGTTGGTAAAC  
AAGGTCAAGCTTTAATTGATCAACTTCGCGCTAATGGTACTCAAACITATAAAGCTACTGTTAAAGTTTACGGAATAAAGA  
CGGTAAGCTGACTTGACTAATCTAGTTGCTACTAAAAATGTAGACATCAACATCAATGGATTAGTTGCTAAAGAAACAGTT  
CAAAAAGCCGTTGACAGACAACGTTAAAGACAGTATCGATGTTCACGACCTACCTAGAAAAAGCCAAGGGTGAAGGTCCA  
TTCACAGCAGGTGTCAACCATGTGATTCCATACGAACTCTTCGAGGTGATGGCATGTTGACTCGTCTCTTGCTCAAGGCATC  
TGACAAGGCACCATGGTTCAGATAACGGCGACGCTAAAAACCCAGCCCTATCTCCACTAGGCGAAAAACGTGAAGACCAAAGG  
TCAATCTCGCAGGTGATGGTATGTTGACTCGTCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTTCAGATAACGGTGACGTA  
GGTACTCAAACCTACAGCGCTACAGTCAATGTCTATGGTAAACAAAGACGGTAAACCAGACTTGGACAACATCGTAGCAACTA  
AAAAAGTCACTATTAACATAAACCGGTTAATTTCTAAAGAAACAGTTCAAAAAGCCGTTGCAGACAACGTTAAAGACAGTAT  
CGATGTTCCAGCAGCCTACCTAGAAAAAGCCAAGGGTGAAGGTCCATTACAGCAGGTGTCAACCATGTGATTCCATACGAA  
CTCTTCGAGGTGATGGTATGTTGACTCGTCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTTCAGATAACGGTGACGTA  
AAACCCAGCCCTATCTCCACTAGGTGAAAACGTGAAGACCAAAGGTCAATACTTCTATCAATTAGCCTTGGACGGAATGTA  
GCTGGCAAAGAAAAACAAGCGCTCATTGACCAAGTTCGAGCAAACGGTACTCAAACCTACAGCGCTACAGTCAATGTCTATG  
GTAACAACAGAGCTGGACAACATCGTAGCAACTAAAAAGTCACTATTAACATAAACCGGTTAATTTCTAA  
AGAAACAGTTCAAAAAGCCGTTGCAGACAACGTTAAGACAGTATCGATGTTCCAGCAGCCTACCTAG

## 4172.5 (SEQ. ID. NO. 301)

ATGAAACTAAAAAGTTATATTTTGGTTGGATATATTATTTCAACCCCTTAACCATTTTGGTTGTTTTTGGGCTGTTCAAAAA  
ATGCTGATTGCGAAAGGCGAGATTTACTTTTTGCTTGGGATGACCATCGTTGCCAGCCTTGTCCGGTGCTGGGATTAGTCTCTT  
TCTCCTATTGCCAGTCTTTACGTGCTTGGGCAAACTCAAGGAGCATGCCAAGCGGGTAGCGGCCAAGGATTTTCTTCAAAATT  
TGGAGGTTCAAGGTCCTGTAGAAATTCAGCAATTAGGGCAAACTTTTAATGAGATGTCCATGATTTCGAGGTAAGCTTTGAT  
TCCTTGAAGAAAGCGAACGAGAAAAAGGGCTTGATGATTGCCAGTTGTTCGCATGATATTAAGACTCCTATCACTTCGATCC  
AAGCGACGGTAGAAGGGATTTTGGATGGGATTATCAAGGAGTCGGAGCAAGCTCATTATCTAGCAACCATTTGGACGCCAGA  
CGGAGAGGCTCAATAAACTGGTTGAGGAGTTGAATTTTTTGACCCTAAACACAGCTAGAAATCAGGTGGAAACTACCAGTAA  
AGACAGATTTTTTCTGGACAAGCTCTTAATTGAGTGATGAGTGAATTTCAAGTTTGTATTGAGCAGGAGAGAAGAGATGTCC  
ACTTGCAGGTAATCCAGAGTCTGCCCGATTGAGGGAGATTATGCTAAGCTTTCTCGTATCTTGGTGAATCTGGTCGATAAC  
GCTTTTAAATATTCTGCTCCAGGAACCAAGCTGGGAAGTGGTGGCTAAGCTGGAGAAGGACCAAGCTTTCAATCAGTGTGACCG  
ATGAAGGGCAGGGTATTGCCCCAGAGGATTGGAAAAATTTTTCAAAACGCTTTATCTGTCGAAACTTCGCGTAACATGAA  
GACAGGTGGTCAATGAGTATTGAGCTTGCAGTTGCGCGTGAATTGGCCCATCAATTGGGTGGGAAATACAGTCAGCAGCCAG  
TACGGTCTAGGAAGTACCTTTACCCTCGTTCTCAACCTCTCTGGTAGTGAATAAAGCCTAA

## 4172.6 (SEQ. ID. NO. 302)

ATGTTTGGTCAAACGGCTCAACATGGTCTTACGAATAGCCTGAAAGACTTCTGGATTTTTCTGCTGAATATAGGTCCACAATT  
GGCGTTTTTTTGGCAGATGCTCCGCTGTTCCAGATCGGTTGAGCAGGGTACTGGAATCACCCTCGTGAGTTCAATATGATTG  
AGCAGATATCTCGCATTTTGGGATGACTCACTTGGGCAAACTCAAGTGGTCTATCAAGAGTCGATTGACCTTGAGTTGCTG  
GTCAATGCATTAATCATCACTTGTCTATTGACAGACTGGTCTCCACGCCAATCAAAATAACGATAGAAATCGCAGGCAG  
TAGTACATGGTCTTGACCTGCTGAAGGGGCGTAAAGACAAAGAGATTATCGACATAAAAAAGTATGTTTCAGGCAGTTAGAACT  
GGCTAGCACGCAACAAATCTGTCCGATAAATCAGCGAGTGCATCATGGTATACTGGCCTTTGGAGAAATTTCCGACCTGGTC  
CCAGCCAAAAATCTGCCGAACAGGCAAGACTGA

## 4174.1 (SEQ. ID. NO. 303)

ATGGAACATTAGCAACTTATTTTTCAACCTATGGAGGAGCTTTCTTCGCTGCATTGGGAATTGTATTGGCGGTTGGATTAAG  
CGGTATGGGGTCTGCTTATGGAGTTGGTAAGGCTGGGCAATCTGCCGACGCTTACTGAAAGAACAGCCTGAAAAAGTTTGGC  
TCAGCTTTGATATTGCAATTATTGCCCGGAACACAAGGATTATATGGTTTTGTTATTGGAATTTTAAATTTGGTTGCAATTAAC  
CCAGAACTTCCTTTAGAAAAAGCGGTTGCTTATTTCTTTGTAGCTCTTCCAATTGCTATTGTAGGATACTTTTCAGCTAAGCAT  
CAAGGAAATGTAGCAGTAGCGGGAATGCAAACTTGGCTAAAAGACCAAAAAGAAATTCATGAAGGGAGCAATTTTAGCTGCC  
ATGGTAGAAACCTATGCAATTCTTGCTTTTGTGCTATCATTCATTTTGACCCTTCGTGTATAA

## 4175.2 (SEQ. ID. NO. 304)

ATGTTAAAAATCAGAAAAACAATCACGTTATCAAATGTTAAATGAAGAATTGTCTTCTATTGGAAGGCGAAACCAATGTTT  
TGGCTAATCTTTTCAAACGCCAGTGCTCTCATAAAATCACGTTTTCTAATACCGTATTTGCAGGCTTTTATTTGTTTCGATGGAA  
AGGAATTTGGTTTTAGGGCCCTTCCAAGGAGGTGTTTCTCGATCCGTATTGCACTAGGCAAGGGTGTGTTGTTGGTGAGGCAGCT  
CACTTTCAGGAAACTGTTATTGTTGGAGATGTGACGACCTATCTCAACTATATTTCTTGTGATAGTCTAGCTAAAAAGTGAAT  
TGTGGTGCCGATGATGAAGAATGGTCAGTTACTTGGAGTTCTGGATCTGGATTCTTCAGAGATTGAGGATTACGATGCTATGG  
ATCGAGATTATTTGGAACAATTTGTCGCTATTTTGCTTGA AAAAGACAGCATGGGACTTTACGATGTTTGAGGAAAAATCTTAA

## 4175.3 (SEQ. ID. NO. 305)

ATGTCAGTATTAGAGATCAAAGATCTTCACGTTGAGATTGAAGGAAAAAGAAATTTTAAAGGGGTTAACCTGACCCTGAAAA  
CAGGAGAAATTGCCGCTATCATGGGACCAAAATGGTACAGGTAATTCGACTCTTTCTGCCGCTATCATGGGAAATCCAAACTA  
TGAAGATACTAAAGGTGAAGTTTGTGTTGATGGCGTAAACATCAGTCTTGGAGTGGATGAGCGTCCGCTATGGGACTT  
TTCCTTGCTATGCAATACCCATCAGAAATCCCTGGAATTAACCAATGCTGAGTTTCTCGTGCCGCTATGAAATGCGGGTAAAGA  
AGATGATGAGAAGATTTTCAAGTTCGTGAGTTTATTACTAAGCTAGATGAAAAAATGGAATTGCTCAACATGAAAGAGAAATG  
GCAGAGCGTTACCTCAACGAAGGCTTCTCTGGTGGTGAGAAAAACGCAATGAAATTTCTCAACTTTTGTGTTGGAGCCAA  
CATTTGCTCTTTTGGACGAGATTGACTCAGGCTTGATTTGACGCTCTTAAAGTTGTGTCTAAAGGTTGCAATGCCATGCCG  
GGTGAAGGTTTTGGTGCTATGATCATCACTCACTACCAACGCTCTTTGAACTATATCACACCTGATGTGGTACAGTGATGAT

GGAAGGTCGTGTTGTCCTTTCTGGTGGTCCAGAATTGGCTGCGCGTTTGGAACTGAAGGATACGCAAAATTAGCTGAAGAA  
CTTGGCTACGACTACAAGGAAGAATTGTAA

## 4174.4 (SEQ. ID. NO. 306)

ATGCCCTACAAAAGACAAAAGGAGTTTCAATGGCACTTTCTAACTAGATAGCCTTTATATGGCAGTGGTAGCAGACCATT  
CGAAAAATCCACATCACAAAGGGAAGTTAGAAGATGCTGAGCAAAATCAGTCTCAACAATCCGACTTGTGGGGATGTCATCA  
ACCTCTCTGTCAAGTTTGTATGCAGAGGACCGTTTGGAAAGATATTGCTTTTCTAAATTCAGGATGCACGATTTCAACTGCTTCT  
GCTAGTATGATGACAGATGCCGTTTATAGGAAAAACCAACAAGAAATTTAGAACTGGCGACTATTTTTCTGAAATGGTTC  
AAGGGCAAAAAGATGAGCGTCAAGACCAACTTGGAGACGCGGCATTCTTGTACGGTGTGGCCAAATTCCTCAAAGAATCA  
AGTGTGCAACCCCTAGCTTGGAAATGCCCTTAAGAAAAACAATTGAAAATCAAGAAAAACAGTAA

## 4175.5 (SEQ. ID. NO. 307)

ATGAAAATTCAGACCTATTGAGAAAAGATGTCATGTTGCTAGATTTCAGGCAACTGAAAAACAGCTGTCATCGACGAGA  
TGATTAATAAATTTGACAGACCACGGTTATGTAACAGATTGTGAAACATTTAAAGAAAGGAATTTTGGCGCGTGAAGCTTTGAC  
TTCTACTGGTTTGGGTGATGGAATCGCAATGCCTCACAGCAAAAACGCTGCTGTCAAAGAAGCGACAGTTCTATTTGCTAAG  
TCAAATAAGGGTGTGACTACGAGAGCTTGGATGGACAAGCAACTGACCTCTTCTCATGATTGCAGCTCCAGAAGGTGCCA  
ATGATACTCACTTGGCAGCCTTGGCAGAATTGTCTCAATACTTGTATGAAAGACGGTTTTCAGACACAACTTCGTCAGCAAC  
ATCTGCAGACCAAGTTATCGAACTTTTGGACCAAGCTTCAGAAAAAACTGAGGAACCTTGTCAAGCACCTGCTAATGACTCT  
GGTGACTTTATCGTAGCTGTACAGCTTGTACAAACAGGTATTGCCACACTTACATGGCCCAAGAAAGCCCTTCAAAAAGTAG  
CTGCTGAAATGGGGTGTGATCAAGGTGCAAAACCAACGGTGTACGGGTGTTGGAATCAACTAACTGCAGAAAGATATCCG  
TAAGGCTAAAGCTATTATCAATTGCAGCAGACAAGGCCGTTGAAATGGATCGATTGATGGAAAACCACTTGATCAATCGTCCA  
GTTGCTGACGGTATCCGTAAGACAGAAGAGCTAATTAACCTGGCTCTTTCAGGAGATACTGAAGTCTACCGTGCCGCTAATG  
GTGCCAAAGCTGCAACAGCCTCTAACGAAAAACAAAGCCTTGGTGGTGCCTTGTACAAACACTTGTATGAGTGGTGTATCTCA  
AATGTTACCATTCGTTATCGGTGGTGTATCATGATTGCCCTTGCCTTCTTGATTGACGGTGCCTTGGGTGTTCCAAATGAAAA  
CCTTGGCAATCTTGGTTCTTACCATGAGTTAGCTTCTATGTTTCATGAAAAATGGTGGAGCTGCCTTGGTGTGATGCTTCCAGT  
CTTTGCCGGTTATGTTGCCCTACTCTATTGCTGAAAAACCGGGTGTGGTAGCAGGTTTCGTGGCTGGTGTATGCCAAAGAAG  
GTTTTGCCCTTGGTAAAAATTCCTTATGCCGAGGTGGTGAAGCAACTTCAACTCTTGCAGGTGTCTCATCTGGTTCTCCTAGGTG  
CCCTTGTGGTGGTATCTCGCAGGTGCCTTGGTTCTTGCATCAAGAAATACGTTAAAGTTCTCTCTCACTCGAAGGTGCT  
AAATCAATCCTTCTATTGCCACTTCTTGAACAATCTTGACAGGATTGTTATGCTAGCTGTGAATATCCCAATGGCTGCAAT  
CAACACTGCTATGAATGACTTCTAGGCGGTCTTGGAGGAGGTTTCACTGCTCTTGGTATCGTCTTGGTGAATGATGG  
CTGTTGACATGGGTGGACAGTTAATAAGCAGCTTATGCTTGGTACAGGTACGCTTGCAGCAACTGTTTCTTCAAGGTGGT  
TCTGTAGCCATGGCAGCAGTTATGGCTGGAGGAATGGTGCACCACTTGAACCTTGTGTCGCAACTCTTCTTTCAGGATAA  
ATTTACTAAGGAAGAAGCTAATCTGCTTTGACAAACATCATCATGGGCTTGTCAATTTATCACTGAGGGAGCGATTCCATTG  
GTGCCGTGACCCAGCTCGTGGATTCCAAGCTTATCCTTGGTTCAGCAGTAGCAGGTGGACTCGTTGGTCTTACTGGTATC  
AAACTCATGGCGCCACACGAGGAATCTCGTTATCGCCCTTACTTCAAATGCTCTCCTTACCTCGTTTCTGTCTTGGTAGGA  
GCAATCGTAAGTGGTGTGGTTTATGGTTACCTACGCAAAACCAAGCATAA

## 4175.6 (SEQ. ID. NO. 308)

ATGGCAACAAGAATACAAGTACAACAAGACGGAGACCGTCTAAAGCAGAAGTGGAAAGAAAAGAAGCGATTCAACGAAT  
GTTGATTTCGTTAGGAATTCGCGATTTTATGATTTTCGCGACCTTCAAATAGGGCTGCAGGTATAACCTTTATAATTTAAT  
TCGCTTGCTAGTGGGTAGCCTAGCTTATCTGGCGATATTCGGCTATTAATCTATCTCTTCTTTTCAAGTGGATACGAAAAACA  
GGAAGGACTCTTATCTGGCTTTTTCACCATATTTGCTGGCTTACTCTTGATTTTGGAGGCCTACTTGGTTGGAAATATGGTTT  
GGACAAGTCCGTTCTAAAAGGGACCATGGCTCAGGTTGTGACAGATCTGACTGGTTTTCGAACGACTAGCTTTGCTGGAGGG  
GGCTTGATCGGGGTCGCTCTTATATTTCCACAGCCTTCTCTTTTCAAATATCGGAACCTTACTTTATTGGTTCTATCTTGATT  
TAGTGGGTTCTCTCCTAGTCAGCCCTGGTCTGTTTACGATATTGCTGAATTTTTCAGTAGAGGCTTTGCCAAATGGTGGGA  
GGGACGAGCGTCGAAAGAGGAACGCTTTGTCAAACAAGAAAGGCTCGCCAAAAGGCTGAGAAAAGGTTGAGATT  
AGAACAAGAAGAGACTGAAAAAGCCTTACTCGATTGCTCTCTGTTGATATGGAACGGGTGAAATCTGACAGAGGAAGC  
TGTTCAAAATCTTCCACCTATTCCAGAAGAAAAGTGGGTGGAACAGAAATCATCTGCTCAAGCTGAACCTTAAATTCCT  
GAACAGGAAGATGACTCAGATGACGAAGATGTTCAAGTTCGATTTTTCAGCCAAAGAAGCCCTTGAATACAAACTTCCAAAGCT  
TACAACCTTTTGCACCAGATAAAACCAAAAGATCAGTCTAAAGAGAAGAAAATTTGTCAGAGAAAATATCAAAATCTTAGAAG  
CAACCTTTGCTAGCTTTGGTATTAAGGTAACAGTTGAACGGGCGGAAATTTGGGCCATCAGTGACCAAGTATGAAGTCAAGCC  
GGCTGTTGGTGTAAAGGTCACCGCATTTCCAATCTATCAGATGACCTCGCTTAGCCTTGGCTGCCAAAGATGTCGGGATTG  
AAGCAAAATCCCTGGGAAATCCCTAATCGGAATTTGAAGTGCCCACTCCGATATTGCCACTGTATCTTTCCGAGCACTATG  
GGAACAATCGCAAACGAAAGCAGAAAATTTCTTGAAATTCCTTATGGGAAGGCTGTTAATGGAACCGCAAGAGCTTTTGAC  
CTTTCTAAAATGCCCACTTGTAGTTGCAGGTTCAACGGGTTACAGGGAAGTCAGTAGCAGTTAACGGCATTATTGCTAGCAT  
TCTCATGAAGGCGAGACCAGATCAAGTTAAATTTATGATGGTTCGATCCCAAGATGGTTGAGTTATCTGTTTACAATGATATTC  
CCCACCTCTTGATTCCAGTCGTGACCAATCCACGCAAAAGCCAGCAAGGCTCTGCAAAAAGGTTGTGGATGAAATGGAAAACCG  
TTATGAACTCTTTGCCAAGGTGGGAGTTCGGAATATTGCAGGTTTAAATGCCAAGGTAGAAGAGTTCAATTCACAGTCTGAGT  
ACAAGCAAAATCCGCTACCATTCATTGTCGTGATTGTGGATGAGTTGGCTGACCTCATGATGGTGGCCAGCAAGGAAGTGG  
AGATGCTATCATCCGCTTTGGGCAGAAGGCGCGTGTGCAGGTATCCACATGATTCTTGGCAACTCAGCGTCCATCTGTTGATG  
TCATCTCTGGTTTGATTAAGGCCAATGTTCCATCTCGTGTAGCATTGCGGTTTCATCAGGAACAGACTCCCGTACGATTTTGG  
ATGAAAATGGAGCAGAAAAACTTCTTGGTTCGAGGAGACATGCTCTTAAACCGATTGATGAAAAATCATCCAGTTCGCTCCCA  
AGGCTCTTTATCTCGGATGACGATGTTGAGCGCATGTGAACTTCAAGACTCAGGCAGATGCAAGTACGATGAGAGT  
TTTGATCCAGGTGAGGTTTCTGAAAATGAAGGAGAATTTTCGGATGGAGATGCTGGTGGTGTATCCGCTTTTGAAGAAGCTA  
AGTCTTTGGTTATCGAAACACAGAAAGCCAGTGCCTTATGATTACGCGTCTTATCAGTTGGATTTAACCGTGCGACCCGT  
CTCATGGAAGAAGTGGAGATAGCAGGTGTCATCGGTCCAGCTGAAGGTACCAAACCTCGAAAAGTGTACAACAATAA

## 4176.1 (SEQ. ID. NO. 309)

ATGAGTTATTTAAAAAATATAAATTCGATAAATCCCACTTCAAACCTTGGTATGCGAACCTTTAAAAACAGGTATTGCTGTTTT  
TCTAGTTCTGTGATTTTGGCTTTTGGCTGGAAAGGCTTCAAATTTGGTGTGTTGACAGCCGTTTTAGCCTGAGGGAGAG  
TTTTGATGAGAGTGTTTCAATTTGGGACTTCGCGTATTCTAGGAAATAGTATCGGTGGACTCTATGCCTTGGTCTTCTTCTTATT  
AAATACCTTTTCCACGAAGCCTTTTGGGTGACCTTGGTAGTTGTTCCAATCTGCACCATGTTAAACATTATGACAAATGTAG  
CCATGAATAACAAAGCAGGGGTTATTGGTGGTGTAGCAGCTATGTTAATCATTACCTATCAATTCCAAGTGGTGAGACAAT  
TTTGTACGTGTTTGTGCGTGTATTAGAAACCTTTATGGGAGTTTGTGCGCAATTATCGTAAATTACGATATTGATCGTATTTCG  
TCTCTTTTATAGAGAAAAAGAAAAATAA

## 4178.2 (SEQ. ID. NO. 310)

ATGAATAAATCAGAAACCGCCACCAACTTATACGCGCTCTTATCACAAAAACAAGATTCATACACAGGCTGAGTTGCAAG  
CCCTTCTTGCTGAGAACGACATTCAAGTAACCCAGGCAACCCCTCTCACGGACATCAAAAAATATGAACCTATCAAAAGTCCG  
CGAAGAAGATAGCGCTTATTATGTTCTTAACAATGGTTCATCTCAAAATGGGAAAAACGCTCTCGAAGCTCTACATGGAAGAC  
GCCCTTGCTGGATGCGCCAGTTCAACACCAAGTCTACTAAAAACCCCTTCTGGACTGGCTCAATCCCTTGGTTCTATCAT  
GATACTTTGAGCTTCCCTGACGCTATCGTACCCCTTTGTGGTAATGATGTCTGTCTTATCATCTGTGAAGATGCAGATACTGCT  
CAAAAGTGCTTTGAAGAAGTGAAGAAAAATTCGCCCCACCATTTTTCTTTGAAGAATAA

## 4179.1 (SEQ. ID. NO. 311)

ATGAAAAGTATAAAATTAATGCTCTATCTTACATGGGAATTCGTGTCTTGAATATTATTTTTCCCATCCTAACTGGAACCTA  
TGTCGCGCGTGTCTTGGACCGAACTGACTATGGTTACTTCAACTCAGTCGACACTATTTTGTCAATTTTTCTTGGCCTTTGCAAC  
TTATGGTGTCTATAAATACGGTTTAAAGGGCTATCAGTAATGTCAAGGATAACAAAAAAGATCTTAACAGAACCTTTTCTAGTC  
TTTTTTATTTGTGCATCGCTTGTACGATTTTGACCACTGCTGTCTATATCCTAGCCTATCCTCTCTTCTTTACTGATAATCCAAT  
CGTCAAAAAAGGTCTACCTTGTATGGGGATTCAACTCATTGCCAGATTTTTCAATCGAATGGGTCAATGAAGCTCTGGA  
ATTACAGTTTTCTCTTTACAAAACTGCCTTCATCCGTATCCTGATGCTGGTCTCTATTTCTTATTTGTTAAAAATGAACACG  
ATATTGTGTCTATACACTTGTGATGAGTTTATCGACGCTGATTAACTACCTGATTAGTTATTTTGGATTAAAAAGAGACATCA  
AACTTGTAAAAATTCACCTAAGTGATTTTAAACCACTCTTCTCCCTCTGACAGCCATGTTAGTCTTTGCCAATGCCAATATGC  
TCTTCACTTTTTAGATCGCCTCTTCTCGTTAAAAACAGGGATGATGTCAACGTTAGTTACTATACCATAGCTCAGCGAATTG  
TGACCGTTATAGCTGGGGTTGTAACAGGTGCAATTGGAGTGAGTGTCCCTCGTCTCAGTTACTATCTGGGAAAGGAGACAA  
AGAAGCCTATGTTTCTCTGGTTAATAGAGGTAGTGAATCTTTAACTTCTTTATCATTCCACTGAGTTTGGACTCATGGTTTT  
AGGACCAAAATGCCATCCTACTTTACGGTAGTGAAGAAATATATCGGAGGCGGCATCTTGACCTCTCTCTTCCGCTTTTCGTACGA  
TTATCTGGCCTTAGATACCAATCTTGGTTCCCAAATTTCTCTTTACCAATGGCTATGAAAAACGTATCACAGTCTATACAGTCT  
TTGCTGGGCTACTGAATTTGGGCTTGAATAGTCTCTCTTTTCAACCATATCGTGGCTCTGAATACTACTTACTGACAACTA  
TGCTATCAGAGACTTCTCTACTTGTCTTCTATATCATTTTCATCCATAGAAAACAACTCATCCACTTGGGACATATCTTTAGCT  
ATACTGTTGATCTCTCTCTTTTCACTTTCCCTTTGTAGCAATTTATTTCCGTGATTAATTTCTGTATCCTGTAGATATGGTCAT  
TAATTTGCCATTTTGATTA  
ATACTGGTTTGATTGTCTTGCTATCAGCTATCTCTTATATTAGTCTACTTGTCTTCACAAAAGATAGCATTTTCTATGAATTTTT  
AAACCATGCTTAGCCTTAAAAATAAATTTAAAAATCATAG

## 4179.2 (SEQ. ID. NO. 312)

ATGAAACAATAACCGTTGAAGATGCCAAACAAATTTGAATTAGAAATTTTGGATTATATTGATACTCTCTGTAAGGACACA  
ATATCAACTATATTATTAACACGGTACTCTGATTGGGGCGGTTGACATGAGGGCTTTATCCCTTGGGACGACGATATTGAT  
CTGTCATGCCTAGAGAAGACTACCAACGATTTATTAACATTTTCAAAAGGAAAAAGCAAGTATAAGCTCCTATCTCTAG  
AAACTGATAAGAACTACTTTAACAACCTTTATCAACACCGACGACTAGCAATAAATTTGATAGCTCGAAATACAAAAAC  
CTATGAGTCTGGTATCTTTATCGATATTTCCCTATAGATCGCTTTGATGATCCTAAGGTCATTGATACCTGTTATAAACTGGA  
AAGCTTCAAACTGCTGTCTTTTCAAGTAAACATAAAAAATTTGCTATAAGGATAGCCTTTTAAAAAGATTGGATACGAACAGCCT  
TCTGGTTACTCTCTCGACCGGTTTCTCTCGTTATTTTGGCAAATAAAATTCGAGAAAGAAATTCAAAAATATAGTCGTGAAAA  
GGGCAATATATGGCTTTTATCCCTTCAAAATTTAAGGAAAGGAAGTCTTCCCAAGTGGTACCTTTGATAAAACAATCGATTT  
ACCCTTTGAGAATTTAAGCCTTCTGACCTGAAAAATTTGATACTATTTTGACACAATTTTATGGAGATTATATGACCCTAC  
CACCAGAAGAAAAACGCTTCTACAGTCATGAATTTACGCTTATAAATTTGGAGGATTAG

## 4179.3 (SEQ. ID. NO. 313)

ATGATAAAAAATCAATCACTAACCATCACACAAAAACAAAGATTTACGAGATCTTGTATCTGACCTAACCATGACCATCCAAG  
ACGGGGAAAAAGGTTGCTATTATTGGTGAAGAAGGAAATGGCAAATCAACCTTACTTAAAAATTTAATGGGGGAAGCTTTGTG  
TGATTTCACTATCAAGGGAAACATCCAATCTGACTATCAGTCACTGGCCTACATTCCCTCAAAAAAGTCCCTGAGGACCTAAAA  
AAGAAAACTTTACACGACTACTTCTTTTAGATTCTATTGATTTAGACTACAGTATCCTCTATCGTTTGGCGGAGGAATTCGAT  
TTTGATAGCAATCGTTTCGAAAGTGACCAAGAGATTGGCAATCTATCAGGGGGCGAAGCTTTGAAAAATTCAGCTTATCCATG  
AGTTAGCCAAACCCCTTTGAGATTCTATTTTAGATGAACCTTCAAAATGACCTAGACCTTGAGACAGTTGATTGGCTAAAAGGC  
CAGATTCAAAAGACCAGGCAAAACCGTTATTTTCATTTCCCATGATGAAGACTTTCTTCTGAAACGGCAGACACTATTGTTCA  
CTTGGCACTGGTCAAAACCCGTAAAGAAGCGGAAACGCTAGTAGAGCATTTAGACTATGATAGCTATAGTGAGCAGAGAAAA  
GGCTAAATTTGCCAAACAAAGTCAGCAAGCTGTCAACAAACCAAGGACCTACGATAAAACCATGGAACCAATCGGAGAGT  
TAAGCAAAATGTAGAAACTGCGCTTCGAGCTACCAAGATAGTACTGCCGGTCCCTATTGGCTAAAAAGATGAAAACTGTC  
CTCTCACAAGAAAAACGCTACGAAAAAGGACGCTCAGTCCATGACTCAAAAGCCACTTGAAGAGGAACAAATCCAACCTTTCT  
TTTCAGACATCCAACCATACCAGCTTCTAAAGTCTTAGTCCAAGTGAAGAAAGAAAAATTTGTCCATTGACGACCGAGTTTG  
GTTCAAAAAACTACAATAACTGTCCGTGGCAAGAAAAAATTCGGTATTATTCGG  
GCCAAATGGTGTGGGAAATCAACTCTGTTAGCCAAGTTACAGAGACTTCTGAATGATAAAAGAGAGATTTCACTTGGTTTT  
ATGCCACAAAGATTACCACAAAAAAGTCAATTTGATTATCCCCAATAGCCTATCTCAGTAAAACTGGGGAAAAAGAGGAA  
CTACAGAAAAATCCAATCTCACCTAGCTAGTCTCAATTTAGTTATCCAGAAAAATGCAGCATCAAAATTCGTCCTTATCTGGCGG  
ACAACAGGGAAAACTCCTGCTTTTGGATTTAGTCTGCGCAACCAAACTTTCTCTGCTGGATGAACCCACACGAACTTTT  
CTCCCACTTCTCAACCCCAATCAGAAAACTCTTTGTACCTATCCAGGCGGTCTCATCACTGTTTCGATGACCGTCTGTTCT  
TAAAGAAGTCTGCTCGATCATCTATCGCATGACAGAACACGGTTTGAAGCTAGTTAATTTAGAAGATTTATAA

## 4179.4 (SEQ. ID. NO. 314)

ATGAAACCAAAACATTTTACAACCTTGCTTGGCGAGCAGAATCTTCCACTTTCCGACCAGCAAAAAAGAACAAATTTGAACGTT  
ATTTTGAGCTTTGGTTCGAGTGAAGATTAATTTGACGGCGATTACGGACAAGGAAGAAGTTTATCTCAACATTT  
TTACGATTTCGATTGCACCCATTCTTCAAGGTTTGTATTTCCCAATGAACTATCAAACTTCTTGATATCGGGGCTGGGGCAGGAT  
TTCTAGTCTACCAATGAAAATTTCTATCCGGAGTTAGATGTGACCATTATTGATTCACTCAATAAGCGCATCAACTTCTTA  
CAACTCTTGGCTCAAGAACTGGATTTGAACGGAGTTCAATTTCTACCACGGACGTGCCGAAGATTTTGCCCAAGACAAGA  
TCCGCTCAATATGATTTTGTAAACAGCTCGTGGGTTGCCGTATGCAAGGTCCTATCTGAATTGACTATCCCTACCTTAAGG  
TTGGTGGCAAACTATTAGCACTCAAGGCTAGCAATGCGCCTGAGGAATTATTAGAAGCTAAGAATGCCCTCAATCTCCTTTTT  
AGTAAGGTCGAAGACAATCTCAGCTACGCGCTACCGAATAGAGATCCGCGCTATATCACAGTGGTAGAAAAGAAAAAGAAA  
CACCATAAATATCCACGTAAGGCTGGTATGCCAAATAAACGCCCACTTTAA

## 4179.6 (SEQ. ID. NO. 315)



ATGAGTATTAACCTAATTGCCGTTGATATCGACGGAACCCCTTGTCACAGCCAAAAGGAAATCACTCCTGAAGTTTTTCTGCG  
CATCCAAGATGCCAAAGAAAGCTGGTGTCAAAGTCGTGATTGCAACTGGCCGCCCTATCGCAGGCGTTGCCAACTTCTAGAC  
GACTTGCAGTTGAGAGACGAGGGGGACTATGTGTAACCTTCAACGGTGCCCTTGTCAGAAACTGCTACAGGACATGAGA  
TTATCAGCGAATCCTTGACTTATGAGGATTATCTAGATATGGAATTCTCTCAGTCGCAAGCTCGGTGTCCACATGCATGCCATT  
ACCAAGGACGGTATCTATCTGCAAAATCGCAATATCGGAAATACACTGTACAGAAATCAACCTCGTCAGCATGCCTATCT  
TCTACCGTACCCCTGAAGAAATGGCTGGCAAGAAATTTGTTAAATGTATGTTTATCGATGAACCAAGAAATTTCTCGATGCTGC  
GATTGAAAAAATTCAGCAGAAATTTACGAGCGCTACTCCATCAACAAATCTGCTCCTTTCTACCTCGAACTCCTTAAAAAGA  
ATGTAGACAAGGGTTCAGCCATTACTCACTTGGCTGAAAAACTCGGATTGACCAAGATGAAACCATGGCAATCGGTGATGA  
AGAAAAATGACCGTGCCATGCTGGAAGTCGTTGGAACCCCGTGTGTCATGGAATGGAATCCAGAAATCAAAAAATCGC  
CAATACATCACCAAAACAAATGACGAATCCGGCGTTGCCATGCCATCCGAACATGGGTACTGTAA

## 4179.7 (SEQ. ID. NO. 316)

ATGACTTGGATTATTCTTGGAGTTATCGCTCTTATTGTTATTTTTGTGATTGTTAGCTATAACGGTTTGGTTAAAAATCGTATG  
CAAACCAAGGAGGCTTGGAGTCAGATTGATGTTTCACTTGAACGTCGCAATGACCTCTTGCCAACTTGATTGAGACTGTAA  
AAGGTTATGCCAAATATGAAGGTTCTACCTTGAAAAGGTGGCAGAACTACGTAACCAAGTGCGCGGCAGCGACTTACCAGC  
AGAAGCTATGAAAGCCAGGTGATGCCCTCACTCGTCAGGTTTTCAGGTTATTTTGCAGTTGCAGAAAGTATCCAGATTGAAA  
GCTAGTGCTAACTTTGTTAAATTTGCAAGAGGAGTTGACAAAACAGAAAAATAAAATTTCTTACTCTCGTCAACTCTATAACA  
GTGTTGTGACGAACTACAATGTAAATAGAACTTTCCCGAGCAATATTATCGCTGGAATGTTTGGATTAAAGCGGCAGAT  
TTTCTTCAAACACCTGAAGAGGAAAAAGTCGGTTCCTAAAGTTGATTTAGCGGTTTAGGTGACTAA

## 4179.8 (SEQ. ID. NO. 317)

ATGTTGTTTGATCAAAATGCAAGCAATAAACGAAAAACCTGGATTGTTGCTGGTATTTTCTACTCTTAGCTCTTGTGGT  
TATGCGGTTGGTTATCTCTTTATAAGATCTGGACTTGGTGGTTTGGTTATGCACTGATTATCGGCTTTATCTACGCTTTGTCT  
ATGATTTTCAATCGACAGAGATTGTCTATGTCCATGAATGGAGCGCGTGAGGTGGATGAGCAACCGCACAGACCTTACC  
ATGTAGTGGAAGATATGGCTCTGGTCTGCTCAGATTCTATGCCCGGTGTTTTCATCATTTGATGATCCAGCCTTAAATGCCTTTG  
CGACAGGTTCTAATCCTCAAAATGCGGCTGTTGCTGCGCACTTCAAGTCTACTAGCTATCATGAATCGTGAAGAACTAGAAGC  
TGTTATGGGACATGAAGTCAGTCATATTCGTAATTATGATATCCGTATTTTCGACTATTGCACTTGCCTTGTCTAGTGCTATCAC  
CATGCTTTCTAGTATGGCAGGTCGTATGATGTGGTGGGTGGAGCAGGTCGCAGACGAAGTATGATGACCGAGATGGAAAT  
GGTCTTGAAATCATTATGCTAGTGGTTTCCCTACTAGCTATTGTACTGGCACCTCTCGCTGCAACCTTGGTTTCAGCTCGCTATT  
TCTCGTCAGAGGGAATTTCTGGCAGATGCATCTAGTGTGAGCTGACTCGCAATCCCCAGGGAATGATTAATGCCCTAGATA  
AGTTGGCAATAGCAAACTATGAGTCGCCACCTGATGCTAGCAGTGGCCTTTATATCAATGATCCTAAGAAAGGTGG  
GGGTTCCAAAACTCTTTTATACCCACCCACCTATCTCAGAACGGATTGAACGTTTAAACAGATGTAA

## 4179.9 (SEQ. ID. NO. 318)

ATGAAATTAATATTCAAGAAATTCGTAAGCAGTCTGAAGGTTTGAACCTTTGAACAAACGTTAGATTTAGTTGATGACCTGC  
GTGCACGTAATCAAGAAATTTAGATGTAAAAGATATCCTTGCAAGTTGGGAAAGTACAATATGAAGACCGTATGTATTTCTT  
AGATTATCAACTATCTTATACCATTTGTTCTTGTCTCGAGTCGAGTATGGAGCCAGTTGAGTTAGTTGAATCTTATCCAGTCAC  
GGAAATTTTTCATGGAAGGGCGCAACTAACCAAGTATGATCAAGAAAGTTTATGATGATGACTTGGTCTTGCCCATCGAAATGGG  
GAGCTTGACCTTGCTGAGAGTGTATCAGACAATATCCTGCTAAACATTCTATCAAGGTCTTGACGGCTGAAGAAGAAGCTG  
GTCAAGGATTTATCTCAGGAAATGACTGGCAAAATCATGACAGAGGAAGAATACCAAGCTCAAAAAGCAGTAAAGAAAGAAG  
AAAACAGTCCTTTTGTGGCTTACAAGGACTATTTGACGGAGATGAATAA

## 4179.12 (SEQ. ID. NO. 319)

ATGGAGTTATTTATGAAAAATCACAACCTATGAAATCTATAAGTTAAAAAAATCAGGTTTGACCAATCAACAGATTTTGAAAG  
TGCTAGAAATACGGTGAAAAATGTTGATCAGGAGCTTTTGTGGGTGATATTGCAGATATCTCAGGTTGCCGTAATCCAGCCGTT  
TTTATGGAACGTTATTTTCAGATAGACGATGCGCATTTGTGCAAAAGAGTTTCAAAAATTTCCATCTTTCTCTATTTTAGATGAC  
TGTTATCCTTGGGATTTGAGTGAAATATATGATGCGCCTGTACTTTTATCAAGGGGAAATCTTGACCTCTGAAATTTCCCG  
AAGGTAGCGGTCGTGGCAGTCGTGCTGTAGCAAAACAGGGAGCTAAGTCAGTTGAAAAAGTCAATCAAGGCTTGGAAAAAT  
GAACTGGTTATTGTCACTGGTCTGGCCAAGGGCATTTGACACAGCAGCTCATATGGCAGCTCTTCAAGATGGCGGAAAAACCA  
TTGCAGTGATTGGAACAGGACTGGATGTGTTTTATCCTAAAGCCAATAAACGCTTGCAAGACTACATCGGCAATGACCATCT  
GGTTCTAAGTGAATATGGACCTGGTGAACAACCTCTGAAATTTTCAATTTTCTGCCCGTAATCGCATCATTTGCTGGACTTTGTC  
GTGGTGTGATTGTAGCAGAGGCTAAGATGCGTTACGTTAGCTGATCTTATTCCTGTGAGCGAGCAATGGAAGAAGGACCGCATGT  
CTTTGCTATTCCTGGTAGCATTTTATAGTGGACTATCAGACGGTTGCCATCATTTGATTCAAGAAGGAGCAAAATTTGGTCACCA  
GTGGGAAGATGTTCTTGGCGAATTTGAATTTAA

## 4181.1 (SEQ. ID. NO. 320)

ATGAAACGTCAATTAGCCTTGGTCTCTTTAGTGGTGGTCAAGATTCAACAACCTGCCTTTTCTGGGTATGCAACACTATGA  
AACAGTCGAAGCTGTACCTTTGCCTACGGCCAACGTCATCACCTCGAAATTCAAATTAAGTATAGAGAAATCGCTAAGGAACAG  
GGCATTCGTCAACCATATCCTCGATATGTCTCTGCTGGGACAAATCACTGCTCAGCCAGACTTTGCGACGATTATATTTTCTTA  
CATTCCTGACAAGCTCTGTGTCGAGTCAAAATCCCTCAAACTATATCTATTTAGCTACCGAAACACGGAGATTTCCACGAAA  
ACTGTATCAACACCATCGGGAAAGACTTGGTCAACTTGCTAGACCTCGCTATTTAGAAAGTCTGGGGAAAAATTCCTCCCGC  
CGGTGGCATTTCAATCGACCCCTACTACAACCTACGGTAAGCAAGGAACCTAAGTATGAGGGCTTGGCAGAACACCGCTCTTC  
CAACACGACCTTTATCCAGAGAAATTTGACAACCGCTAA

## 4181.2 (SEQ. ID. NO. 321)

ATGACCGAAACGGTAGAAGATAAAGTAAGTCATTCAATTACTGGGCTTGATATCCTCAAGGGGATAGTTGCTGCGGGAGCTG  
TCATAAGTGGAACCGTTGCAACTCAACGAAGGTATTTACAATGAGTCAGCAGTACTTGAAAAAACTGTAGAGAAAACGG  
ATGCTTTTGGCAACAAATGATACAGTAGTTCTAGGTACGATATCTACAAGTAATTCAGCGAGTTCAACTAGTTTGTGAGCTTCA  
GAGTCGGCAAGTACATCTGATCTGAGTCAGCCTCAACCCAGCGCTTCGACCTCAGCAAGTACAAGTGATCAGAAATCAGCAA  
GTACATCGGCTTCGACAAGTATTTCTGCAATCATCTACTGTGTTAGGTTTACAAACAGCTGCGGCTACAGAAGCAACTGCTAA  
GAAGGTGCAAGAAAGATCGTAAGAAACCAGCTAGTGATTATGTAGCATCAGTTACAAATGTCAATCTCCAATCTTATGCTAAG  
CGACGCAAGCGTTCAAGTGGATTCCATCGAGCAATTTGCTGGCTTCTATAAAAAATGCTGCTGTTTTTCTGGCAATACGATTGT  
AAATGGCGCCCTGCAATTAATGCAAGTCTAAACATTGCTAAAAAGTGAGACAAAAAGTTATACAGGTGAAGGTGTAGTCTG  
GTATATCGTGTTCATTTACTATAAAATTTGAAAGTGACAAATGATGGTTCAAAATTTGACCTTTACCTATACGGTTACGTATGT  
GAATCCTAAAAACAAATGATCTTGGTAATATATCAAGTATGCGTCTCGGATATTCTATCTATAATTCAGGTACTTCAACACAAA

ATGGGGGTCGAAACCTTGGTTTTATTCTAGCATCTGCTGGCTGGCCATCGGGCTTGGTTCGGTTTGGAAAGTTTCCCTACATGAC  
TGCTGCTAATGGCGGTGGAGGCTTTTTACTAATCTTTCTCATTTCCACTATTTAATCGGTTTCCCTCTCCTGCTGGCTGAGTTT  
GCCCTTGGCCGTAGTGGCTGGCGTTTCCGCTACAAAACCTTTGGAAAACTGGGCAAGAATAACAAGTACAACCTTTATCGGTT  
GGATTGGCGCCTTTGCCCTCTTTATCCTCTTATCTTTTACAGTGTTATCGGAGGATGGATTCTAGTCTATCTAGGTATTAGT  
TTGGGAAATTTGTTCCAACCTTGGTGGAACGGGTGATTATGCTCAGTTATTTACTTCAATCATTTCAAATCCAGCCATTGCCCTA  
GGAGCTCAAGCGGCCCTTATCTCTATTGAATCTTCTATGTCATCAGTGGGGTTCAAAGGGGATTGAAAGAGCTTCGAAAG  
TCATGATGCCCTGCTCTTTATCGTCTTTGTTTTATCATCGGTCGCTCTCAGTTTGGCAAATGCCATGGAAGGGGTTCTTT  
ACTTCTCAAACCAGACTTTTCAAACCTGACTAGCACTGGTCTCCTCTATGCTCTGGGACAATCTTTCTTTGCCCTCTCACTAG  
GGGTACAGTCATGTTTGACCTATGCTTCTTACTAGACAGAAAACCAATCTAGTCCAGTCAGGAATCTCCATCGTAGCCATG  
AATATCTCGATATCCATCATGGCAGGTCTAGCCATTTTCAAAGCTCAGTCCCTCTAATATCGTCTGAAGGGGGACCCAG  
CCTGCTCTTTATCGTCTTGCCTCAACTCTTTGACAAGATGCCTTTTGAACCATTTTCTACGTCCTCTTCTCTTGTCTTCTCTT  
TTTGGCAGATGCTCACTTTTCTGTGCTGATGCTGGAAATCAATGTAGACAATATCACCAACCAAGGATAACAGCAACCGTCCCA  
AATGGAGTGTTATTTAGGAATTTTGACCTTTGCTTTGGCATCTCTTACGCCCTATCTTACGGGTGTCTGGCGGATGTTTACA  
TTTTTGTAAGACCTCTTTGACGCTATGGACTCTTGGTTTCCAATCTCTCATGCCATTTGGAGCTCTCTACCTTTCATCTTT  
TACAGGCTATATCTTTAAAAGGCTCTTGCAATGGAGGAACCTCATCTCGATGAAAGAGCATGGAAACAAGGACTGTTCCAA  
GTCTGGCTCTTCTCTCTCGTTTCTCGTATTCCAATCATCATCATGTGGTCTTCATTGCCCAATTTATGTAATCAA  
AGGACTTGTAGTAG

ATGTTGAAAAAATGCGAGTTAAAAAGATGTTATCTTGCTTGCTTTCTTGCTATCTTTTTTGGTGGGGTTTTTCGTTGGTTCAGGA  
TATGTGTATAATATTCTCAGTCTACTCTTAACACCTCTTGGTTTGCAAGGCCTTTGCCAATGAAATCCTCTTCGGTCTCTGGTGT  
ATGGCTGCGCCCATTTGCTGCCATCTTTGTTCCGAGAGTCGGAAGTGCAACGATTGGAGAAGTGCTAGCTCGCTCTGTCTGAAG  
TCCTTTATGGTAGCCAATTTGGTCTAGGAAGCTCTTTGTCTGGCTTTGTTCACAGGTTTGGGAAGTGAATTTGGTTTATCATGTA  
CTAAGAATCGCTATGAAAGTTGGCTCTCTCTAACTGCTAATAGTATTGGGATTACGCTTGTTAGCTTTGTCTATGAATACATT  
AAGTTAGGTTACTACGCCCTTTTCCCTTCCGTTTGCTTTCTCTTGCTTTGCTTTGCTTGTTGTTAGCTTTATTTCTGTTTATTCTCTGTACCATT  
CTTGGTTTCGTGCCAATTGTCAAACCTCTATCATCAGTTTGCAACTGGAGGAAAGCATAG

ATGGTCAAAGTAGCAACCCAGACACCGATTATCAGTCTCTTCTTGCTGATTTTATCCTTGAAACATCTTTCATTCTTCGATT  
GCTCTGACTCTTTCGGTAGTCGCAATTTGTAATCTCTTTATGCTCTATTACCGTCGATTTAAAAATGTTAGCTTGGATGATCATAG  
CTTGCCATTTTACCATCTTTTGCCAACACTACCTGGCAGTTCAGTTACACGGAGATGCTTACACAGGCATGCTTGGAAACGAG  
GGCCTTTGTGACAGCTTTGATCTCGGCCTTGCTTTGTTTCTCTGTTTACATAAAAGAGCTTCTCTTGACTGGCTCAAAAGGG  
GCTATCACGCTCTTGGTCTATGCTTGATTGTTGGTATTCAATTCTTTTCTCTCATTCAGCAAGAAATCAAGTCCCTCAAAGA  
AGCTTGCCCTATTACGTGGTCAAGAAGACTACATTTTGGTCGCCCTGATTTACAGAAAGGTTCTGATGACAGTCTTTAGGTGGC  
GCCATCTTTACCTGAGAGCTCTACTGCTACGGGATGACGAAACATGCACAGTTGAAGAAATAGCTATCGGACTTTTATATT  
CCTAAAAAACAATAATCTACCTGCTTTTCTTTTATGTTCTTCAAAACAGCTATTTTTATAA



## 4183.7 (SEQ. ID. NO. 325)

ATGAGAAAGCACCAATTACAAAGTTCACAAATTAACCATTTTATCTATGATGATTGCCCTTGATGTAGTCCTTACACCTATCTT  
 TCGAATTGAGGGAATGGCACCGATGTCCAGTGTAGTCAATATTCTAGCAGGAATCATGATGGGACCTGTTTATGCCTTGGCT  
 ATGGCTACAGTCACAGCCTTTATCCGTATGACGACTCAAGGGATTCCGCCTTTAGCTCTCACAGGAGCGACTTTTGAGGCCCT  
 TCTAGCAGGTCTCTTTTATAAGTACGGTCGAAAAATTTCACTATTCTGCTCTAGGAGAGATTTTGGGAACAGGTATTATTGGTT  
 CCATTGTTTCTATCTGTTATGGTACTCTTTACAGGATCAGCTGCTAAGCTTAGCTGGTTTATCTACACGCCTCGATTTTTCG  
 GAGCAACCTTGATTGGTACAGCGATTTCCTTTATTGCCTTTCGATTTTTAATCAAGCAGGAATTCTTTAAAAAAGTGCAGGGA  
 TATTTCTTTAGTGAAAGGATAGACTGA

## 4183.8 (SEQ. ID. NO. 326)

ATGCAGGAATTTACAAATCCCTTTCTATAGGCTCTAGTTCCCTCATTCACTGCATTACCAATGAGATTTCTTGTGAGATGCTG  
 GCAAATGGGATTTTGGCTCTGGGATGCAAACCTGTCTATGGCAGATGATTCCCGTGAAGTTCTTGATTTTACTAAGCAAAGTCA  
 GGCTCTCTTCATCAATTTGGGGCATTGTGTCAGCTGAGAAGGAAAAAGCAATCCGCATGGCAGCTTCGTATGCAAACCAATCT  
 TCTCTCCCGATGGTAGATAGTCGGTTGGCGTAACGACTTCATCCATTTCGTAAGAGCTTAGTTAAAGACCTTTTAGACTATAG  
 ACCTACGGTCCTTAAAGGAAACATGTGAGAAATTCGAAGTCTTGTGGATTAAAGCACCACGGCGTTGGGGTCGATGCGAGT  
 GCTAAAGATCAAGAAACGGAGGATTTCCTTCAAGTCTTGAAGACTGGTGTGACACCTATCCTGGTATGCTTTCTTAGTCAC  
 AGGTCCCAAGGACCTCGTCGTTTCGAAAAATCAGGTCTGCTGACTGGGAAATGGCTGACTGAATTAGACTGGATAACAGGG  
 ACAGGAGACTTGGTTGGAGCCTTAACAGCTGTTTTCTCAGCCAAGGAAAGACTGGTTTTGAAGCTTCTTGCCTAGCAGTCTC  
 TTATCTCAATATCGCTGCTGAGAAAAATAGTTGTTCAAGGAATGGGATTGGAAGAATTTCTGTACCAAGTACTCAATCAGCTTT  
 CGCTCCTAAGAAGAGATGAAAAATGGCTAGATACCATCAAAGGAGAGGTTTATGAATAG

## 4185.3 (SEQ. ID. NO. 327)

ATGAACCATAAAAATCGCAATTTTATCAGATGTTTCATGGCAATGCGACGGCGCTAGAAGCAGTGATTGCAGATGCTAAAAATC  
 AAGGGGCCAGTGAATATTGGCTTCTGGGAGATATTTTCTTCTGGTCCAGGCGCAAATGACTTAGTCGCCCTGCTAAAGGAC  
 CTTCTATCACAGCAAGTGTTCGAGGCAATTGGGATGATCGTGTCTTGAAGCTTTAGATGGGCAATATGGCTTAGAAGACC  
 CACAGGAAGTTCAGCTTTCGGTATGACACAGTATTGATGGAGCGAATGGATCCTGCAACGATTGTCTGGCTACGAAGCTT  
 GCCTTTGCTGGAAAAAGAAAGAAATTTGACGGATTGCGCTTTTCTATCTCTCATAATTTACCTGACAAAAACTATGGTGGTACT  
 TGCTAGTTGAGAATGATACAGAGAAATTTGACCAACTGCTAGATGCGGAAACGGACGTGGCAGTTTATGGTCACTGTTTACAA  
 GCAGTTGCTTCGTTATGGAAGTCAAGGGCAACAAATCATCAATCCAGGGTCGATTGGCATGCCCTATTTTAAATGGGAGGGC  
 TTAAAAAATCACCGTTCCCGATGATGCCGTGATAGAAGTTGAAGATGGGGAATTACTCAATATCCAATTTCTGTAAGTTGCTTA  
 TGATTACGAAGCTGAGTTAGAATTTGGCCAAGTCCAAGGGGCTTCCCTTTATCGAAATGTATGAAGAACTGCGTCGTGACGAT  
 AACTATCAGGGGCACAATCTGGAATTATTAGCCAGCTTAATAGAAAAGCATGGGTATGTAGAGGATGTGAAGAATTTTTTTG  
 ATTTTTGTAA

## 4186.1 (SEQ. ID. NO. 328)

ATGAATGTAAATCAGATTGTACGGATTATTCCTACTTTAAAAAGCTAATAATAGAAAAATTAATGAAACATTTTATATTGAAAC  
 CCTTGGAAATGAAGGCCCTTGTAGAGAATCGGCCCTTCTGTCACTAGGTGACCAACGGGTCTTGAAAAAGCTGGTTTTAGAA  
 GAAGCTCCAGTATGCGTACTCGTAAGGTAGAGGGAAGAAAAAACTAGCTAGATTGATTGTCAAGGTGGAAAAATCCCTTA  
 GAAATTAAGGAATCTTATCTAAAAACAGATTGATTATCGATTATATAAAGGTCAAAATGGTACGCTTTTGAAATTTTCTC  
 ACCAGAAGATGATTGATTTGATTTGATTCATGCGGAAGATGACATAGCAAGTCTAGTAGAAGTAGGAGAAAAAGCCTGAATTTCAA  
 ACAGATTTGGCATCAATTTCTTAAAGTAAATTTGAGATTTCTATGGAATTACATCTCCCACTGATATCGAAAGTTTCTTGA  
 ATCATCTGAAATTTGGGCGATCCCTTGATTTTATTCCAGCTCAGGGGCGAGATTGACTGTGGACAATACGGTTACCTGGGACT  
 TATCTATGCTCAAGTCTTGGTCAATGAATTAGACATAGCAAGTCTTCGCCAGAAGTTTGAGTCTACTGAATATTTTATTCCTA  
 AGTCTGAAAAATTTCTCTTGGTAAAGATAGAAATAATGTTGAATTGTGGTTTGAAGAAGTATGA

## 4186.2 (SEQ. ID. NO. 329)

ATGAAGTGGACCAAGATTATTAAAAAAATAGAAGAACAATCGAGGCAGGGATTTATCCCGGAGCCTCTTTTGGCTATTTTA  
 AGGACAATCAATGGACAGAGTTCTATTTAGGCCAGAGTGACCCAGAGCATGGCTTGCAGACTGAGGCAGGACTAGTTTATGA  
 CCTAGCTAGTGTGACGAAGGTGTTGGGGTTGGCACAGTTTGTACCTTCTTGTGGGAAATAGGTCAATTAGATATTGATAGAC  
 TGGTAATAGATTGTTTACCTGAGAGTGATTATCCAGACATCACTATTCGCCAGCTCTTGACTCATGCAACAGACCTTGATCCT  
 TTTATCTCAATCTTGATCTTTTAAACAGCCCTGAATTAAGGAAGCGATGTTTCACTCAACAGACGAAGTCAAGCCAGCCTT  
 TCTTTATTCGGATGTCATTTTGTGTTGGGCTTTATTTTGGAAAGAATTTTAAATCAAGATTGGATGTGATTTTAAAGGA  
 TCAAGTCTGGAAACCTTGGGGAATGACGGAAACTAAGTTTGGGCCAGTTGAGCTTGCTGTTCCAACAGTTAGAGGTGTAGAG  
 GCAGGCATAGTGCATGATCCCAAGGCTCGTCTCCTGGGTAGACATGCTGGGAGTGTGTTTATTTTCGACTATAAAGGATTT  
 ACAAATCTTTTGAACACTATTTAGCAGATGATTTTGAAGAGACTTAAATCAAAATTTTCTCCTTTGGATGACAAGGAAC  
 GTTCTTTAGCATGGAATTTGGAAGGAGATTGGCTAGACCATACGGGCTATACAGGTACCTTTATCATGTGGAATCGTCAGAA  
 GCAAGAAGCCACTATTTCTATCGAATCGTACCTATGAAAAGGACGAGAGAGCTCAATGGATATTAGACCGCAATCAAGTG  
 ATGAACCTGATTGCAAGAAGAGATGA

## 4187.2 (SEQ. ID. NO. 330)

ATGATGAAGAAGACTTATAATCATATTTGGTCTGGGGAGTCAATTTCTATAGCATTGCAATTGCTGTTTTGCTTTACTCCT  
 CAAGAACAATCTACCGTGGGAGTGGGAACCTCAGGTATTCAGCATCTTGGACGCCTGGTTTTTCTTTGACTCCTTTCAATTC  
 TCTCTGGAACCTGGGCGAAGTGAGTGACATTGGACAATTAATGTTGGATTTTTTACAAAATATCCTCAATGTCTTCTGTTTTT  
 TCCTCTGATTTTCCAACCTCCTTATCTATTTCCAAATTTGGCGAAAAACAAAAAAGGTCCTTCTTTTAGTTTTCTTGTGAGTCTT  
 GGAATCCAGTGTACGCAATTAATCTTGGACTTTTCTTGTGTTTCAATTCGCGTCTTTGAGATTGATGATTGTGGACCAACACT  
 TTGGGTGGCTATCTGGCTTGGCTCCTTTATAACGATTACATAAAAAACAAGGTAAGGAATTA

## 4188.1 (SEQ. ID. NO. 331)

ATGAAGATTCTCTCTTAACTTTTGCAAGGCATAAATTTGTTTATGTCTTGCTTACTTTGCTTTTTCTTGCTTTGGTTTATCGTG  
 ATGTTTTGATGACTTATTTCTTTTTTGATATTCATGCGCCCGATCTAGCTAAATTCGATGGACAAGCAATTAATAATGACTTAT  
 TAAAATCAGCATTAGATTTTCGTATTCTCCAGTTCAATCTAGGTTTTTATCAATCATTTATTATTCCAATCATCATTTGTTTGTCT

AGGTTTTCAATATATTGAGCTGAAAAATAAAGTTTTACGATTGAGTATTGGAAGAGAAGTGAGTTATCAAGGGTTAAAAAGA  
AAGTTGACTTTTGAAGTTGCAAGTATCCCTTGTGGATATATTTAGTGACTGTGCTGATAATTGCAATTATAACCTATTTCTTT  
GGGACTTTTTCTCCTCTTGGATGGAATTTCTATTTTCTGATGGAAGTGGTTTACAAAGACTCCTAGATGGAGAGATAAAAAAG  
CTATTTGTTCTTTACTTGTGCTCTACTAATCGGTATTTTCATCAATGCAATCTATTTTTTACAAATAGTTGATTATGTGGGGAAT  
GTGACTCGTTCGGCAATCACCTATTTGATGTTTCTTTGGCTTGGTTCTATGCTGCTTATAGTGCCTTGCCTTACTATATGGTTC  
CTATGACGAGTTTGTATGCAAGCTAGCTATGGGGATGTAAGTTTGTATGAAACTCTTTACTCCTTATATCCTTTATATTGTCCCTT  
ACATGGTGTGTTGAAAAATATGAAGATAATGTTTAA

## 4188.2 (SEQ. ID. NO. 332)

ATGAAGATAATGTTTAAAGAATTTTAAACAATATTTTGTCTAAATAGAAAGATTGTTTTACTACTTCGTATAGTTCTGATGATGAT  
TTTGATAAACCATCTATTGTCAACAGCGGTTCAAAAGCAGGATGCTGTTATCTTTTTCAAGAGAGAATTGATTTCATTTTTTCT  
CTATAATGACTATTCTGAAGCGAATTTAGAAATCCCCAACTATTGTTAAACCTTTCGCTTTTCATGGTAGGATGGCTCTCTGT  
CATTTTACTTGAAGTGATTTTGGCAGACCATTACCATCACTTGTATCGCTATCAATCAAGCTCCTTTTCGATTATACAAGGA  
AACGATTGGTTGTCTATTTCTAAATTTTTTACTCAAGATTGTTTGTCTGGTTTCTTGGTTTACTTCTCTAGGAATTCATTTCAA  
AACAGTCGCACTTTTCTTTTACTTGTCTAGTTAATGATGTTGTACTTACTACTGCTTATCTGATAGCACTGATTAGTGGGG  
CGCTGGTTTTCTTTCTTCTATTTTTCATTTTTCAGCATTTTGTGGGACAAGAATGGATGATGGATCATATTGTAACAGTGTATTAGT  
ACTCTTAAAGTTTATTAGTTATGTTGATTGTTAGTTCGCTTGAAGAGAAATTTAAGAAAGGATAA

## 4188.5 (SEQ. ID. NO. 333)

ATGGGCAAAGGAGAGATGGGCAAAGGAGTTATTGGCTTGGAGTTCGACTCAGAAGTATTGGTCAACAAGGCTCCAACCTTCT  
AATTGGCAAATGGTAAAACAGCGACTTTCCTAACCAGTATGATAGCAAGACCTTGTGTTTGCAGTAGATAAGGAAGATAT  
CGGACAGGAAATATTGGTATAGCTAAAGGAAGCATCGAAAGTATGCATAATCTTCTGTAAATCTAGCAGGTGCCAGAGTT  
CCTGGCGGAGTAAATGGTAGCAAAAGCAGCGGTGCATGAAGTCCAGAATTTACAGGGGGAGTTAATGTCACAGAGCCAGCT  
GTTTATGAAATCGCAGAGTATAAGGGATCTGATTTCGTTGTAACCTTCTTACTACAAAAAAGATTATCTTACAAAGCTCCTCT  
TGCTCAGCAGGCACTTCTGAAACAGGAAACAAGGAGAGTGACCTCTAGCTTCACTAGGACTAACAGCTTCTTCTCTGGT  
CTGTTTACGCTAGGGAAAAAGAGAGAACAATAA

## 4188.10 (SEQ. ID. NO. 334)

ATGTTTAAAGTTTTACAAAAAGTTGGAAAAGCTTTTATGTTACCTATAGCTATACTTCTGCAGCAGGTCTACTTTTGGGGAT  
TGGTGGTGCACTTTCAAACCCAACACGATAGCAACTATCCAATACTAGACAATAGTATTTTTCAATCAATATTTCCAAGTAA  
TGAGCTCTGCAGGAGAGGTTGTATTCAGTAATTTGTCACTACTTCTCTGTGTTGGGATTATGTATTGGCTTAGCGAAACGAGAT  
AAAGGAACCGCTGCGTTAGCAGGAGTAACTGGTTACTTAGTTATGACTGCAACGATCAAAGCTTTGGTAAAACTTTTATGG  
CAGAAGGATCTGCAATTGACTCGGAGTTATTGGAGCATTAGTTGTGCGGAATAGTTGCCGTATATTTGCACAACCGATATAA  
CAATATTCAATTACCTTCGCTTTAGGATTTCTTTGGAGGTTCAAGCTTCTCTATTGTTTACATCGTTTCTTCTATCTTGATT  
GGCTTTGTCTTCTTTGTTATTTGGCCACCTTTCCAACAACCTTCTTGTCTTACAGGTGGATATATTTCTCAGGCGGGTCCAATT  
GGAACCTTTTCTATATGGATTTTTAATGAGACTTTCTGGAGCAGTAGGCTTACATCATATAATTTACCTATGTTTGGTATACT  
GAACCTGGTGGTGTGAACTGTTGCAGGACAAACAGTGGTTGGAGCTCAAAAAATATTTTTTGTCTAATTAGCCGATTTGGC  
CCATCTGGGATTATTTACAGAAGGAACAAGGTTTTTGGAGGTCGTTTCTCAACAATGATGTTCCGTTTACCGGCTCGCTGTT  
AGCGATGTACCATAGTGTTCCTAAAAATCGTCGTAAAAAATACGCGGGTTTGTTTTTTGGAGTTGCTTTAACATCTTTTATTAC  
CGGTATTACAGAACCAATTGAATTTATGTTTCTATTTCGTCAGTCCGGTTCTATATGTTGTTACCGCATTCCTTGATGGTGTAG  
CTTCTTTATTGCAGACGCTTAAATATTTCAATAGGAAACACATTTTCAGGAGGTGTAATCGATTTCATTTATTTGGAATTTT  
GCAGGGGAACGCTAAGACGAATTGGGTTCTTCAGATTCCATTTGGACTTATTTGGAGTGTTTTGTATTATATTATTTTATAGATG  
GTTTACTTCAATTCAACGTTCTAACGCCAGGGCGAGGAGAAGAAGTAGATTCTAAAGAAATTTCTGAATCCGCAGATTCA  
ACTTCAAATAGTCAGATTATTTAAAACAGGATAGCCTACAAATTATCAGAGCCTTGGGTGGATCAAATAATATAGAAGATG  
TAGATGCTTGTGTGACACGTTTACGTGTAGCTGTAAAAAGAAGTTAATCAAGTTGATAAAGCACTTTTAAAAACAATTTGGTGC  
AGTTGATGTCTTAGAAGTGAAGGGTGGCATTCAAGCAATCTATGGAGCAAAAGCAATCTTATATAAAAAATAGTATTAATGAA  
ATTTTAGGTGTAGATGATTAA

## 4188.11 (SEQ. ID. NO. 335)

ATGAAATTTAGAAAATTAGCTTGTACAGTACTTGCAGGTGCTGCGGTTCTTGGTCTTGTGCTTGTGGCAATTCTGGCGGAAG  
TAAAGATGCTGCCAAATCAGGTGGTGACGGTGCCAAACAGAAATCACTTGGTGGGCATTCCAGTATTTACCCAAGAAAAA  
ACTGGTGACGGTGTGGAACTTATGAAAATCAATCTCGAAGCGTTTGAAGAAAGCAAAACCCAGATATAAAAGTGAAATGT  
GAAACCATCGACTTCAAGTCAGGTCTGAAAAAATCACACAGCCATCGAAGCAGGAACAGCTCCAGACGTACTCTTTGATG  
CACCAGGACGTATCATCCAATACGGTAAAAACGGTAAATGGCTGAGTTGAATGACCTCTTACAGATGAATTTGTTAAAGA  
TGTCACAATGAAAAATCTGTACAAGCAAGTAAAGTGGAGACAAAGGCTTATATGTATCCGATTAGTTCTGCCCATTTCTAC  
ATGGCAATGAACAAGAAAAATGTTAGAAGATGCTGGAGTAGCAACCTTGTAAAAAGAAGGTTGGACAACCTGATGATTTGAA  
AAAGTATTGAAAGCACTTAAAGACAAGGGTTACACACCAGGTTCAATTGTTCAAGTTCTGGTCAAGGGGGAGACCAAGGAACA  
CGTGCCCTTATCTTAACCTTTATAGCGGTTCTGTAAACAGATGAAAAAGTTAGCAAAATATACAACCTGATGATCTTAAATTCGT  
CAAAGGCTTGAAGAAAGCAACTAGCTGGATTAAAGACAATTTGATCAATAATGGTTCACAATTTGACGGTGGGGCAGATATC  
CAAACTTTGCCAACGGTCAAACATCTTACACAATCTTTGGGCACCAGCTCAAAATGGTATCCAAGCTAAACTTTTGAAG  
CAAGTAAGGTAGAAGTGGTAGAAGTACCATTCCCATCAGACGAAGGTAAGCCAGCTCTTGAGTACCTTGTAAAACGGGTTTGC  
AGTATTCAACAATAAAGACGACAAGAAAGTTCGCTGCATCTAAGAAATTCATCCAGTTATCGCAGATGACAAGGAGTGGG  
ACCTAAAGACGTAGTTTCGTACAGGTGCTTTCCAGTCCGTACTTCAATTGGAAAACTTTATGAAGACAACGCATGGAAAAA  
ATCAGCGGCTGGACTCAATACTACTCACCATACTACAACACTATTGATGGATTGCTGAAATGAGAACACTTTGGTTCCCAAT  
GTTGCAATCTGTATCAAAATGGTGACGAAAAACCAGCAGATGCTTTGAAAGCCTTCACTGAAAAAGCGAACGAACAATCAA  
AAAAGCTATGAAACAATAG

## 4188.12 (SEQ. ID. NO. 336)

ATGCAATCTACAGAAAAAACCATTAACAGCCTTTACTGTTATTTCAACAATCATTTTGTCTTGTGACTGTGCTGTTTCATC  
TTCCATTCTACTGGATTTTACAGGGGCATTCAAATCACAACTGATACAATTTGTTATTTCTCCTCAGTGGTTCCCTAAAAATG  
CCAACCATGGAAGCTTCCAACAACCTCATGGTGCAGAACCTGCTTGCATGGATGTGGAACCTCAGTATTTATCTCATTGGT  
AACCATGTTCTTATTGTTGCAACCTCATCTAGCAGGTTATGTTATGGCTAAAAAACGTTTCTATGGTCAACGCATTTCTATT  
TGCTATCTTTATCGCTGATGGCGCTTCCAAAACAGTTGCTGTTGACCAATTGGTACGTATCGTCAACTCATGGGAATCCA  
TGATACTCTCTGGGCAGTTATCTTTGCCCTTTGATTGGATGGCCATTCCGTGCTTCTCATGAAACAGTTTCACTGAAAAATATCCC  
TACAGAGTTGCTTGAATCAGCTAAAAATCGACGGTTGTGGTGAGATTCTGACCTTCTGGAGTGTAGCCTTCCCGATTGTGAAAC

CAGGGTTTGCAGCCCTTGAATCTTTACCTTCATCAATACTTGGAAATGACTACTTCATGCAATTGGTAATGTTGACTTCACGTA  
ACAATTTGACCATCTCATTGGGGTTGCGACCATGCAAGGCTGAAATGGCAACCAACTATGGTTTGATTATGGCAGGAGCTGC  
CCTTGCTGCTGTTCCAATCGTCACAGTCTTCTAGTCTTCCAAAAATCCTTCACACAGGGTATTACTATGGGAGCGGTCAAAG  
GATAA

## 4191.1 (SEQ. ID. NO. 337)

ATGAAAAAAGTAAAGATAATTTTCTACTGGTGTAGGCTGTTTTGCCTTCTCCACTCTCTGTTTTGCCATTGATTTCAAGATAAACT  
CTTATCAAGGGGATTTGTATATTCATGCAGACAATACGGCAGAGTTTAGACAGAAGATAGTTTACCAGTTTGAGGAGGACTT  
TAAGGGCCAAATCGTGGGACTTGGACGTGCTGGTAAGATGCCTAGCGGGTTTGACATTGACCCTCATCCAAAGATTCAGGCC  
GCGAAAAACGGTGCAGAACTAGCAGATGTGACTAGCGAAGTAACAGAAGAAGCGGATGGTTATACTGTGAGAGTCTATAAT  
CCAGGTCAGGAGGGCGACATAGTTGAAGTTGACCTCGTCTGGAACCTTAAAAAATTTACTTTTCCTTTATGATGATATCGCTGA  
ATTAATTTGGCAACCTCTGACAGATAGTTTCAAGAGTCTATTGAAAAGTTTGAATTTTATGTAAGGGGAGACAAGGGGGCTGAA  
AAACTCTTTTCCATACAGGGGAACTTTTATAGAGGGAAGCAAGATGTAAGAGTAACCTTGATTATCTATCCGTTTAGACA  
ATCTTCCGGCTAAGCGTGGAGTTGAGTTGCATGCCTATTGGCTCGGACCGATTTTGCTAGCGCTAGGGATCAGGGATTGAA  
AGGGAATCGTTTGAAGAGTTTAAATAAGATAGAAGACTCGATTGTTAGAGAAAAAGATCAGAGTAAACAACCTCGTTACTTG  
GGTCTCCCTTCGATCTTCCATCTCCTTGTTATTGAGTGTCTGCTTCTATTTTATAGAAGAAAGACCACTCCTTCAGTC  
AAATATGCCAAAAATCATCGTCTCTATGAACCACCAATGGAATTAGAGCCTATGGTTTTATCAGAAGCAGTCTACTCGACCTC  
CTTGGAGGAAGTGAGTCCCTTGGTCAAGGGAGCTGGAATAATCACCTTTGATCAACTTATTCAAGCTACCTTGCTAGATGTGA  
TAGACCGTGGGAATGTCTCTATCATTTTCAAGGAGATGCAGTTGGTTTGGGCTAGTAAAAAGAAAGATGGTTTGTCAAGCTT  
TGAGAAAGACTGCTTAACTAGCTTTTTCAGGTAAAAAAGAAAGAACTCTTTTCCAAATTTGTTTGGCGGATACAAGGTATCTG  
ATAGTCTTTATCGTAGAGCCAAAGTTTCTGATGAAAAACGGATTCAAGCAAGAGGGCTTCAACTCAAATCTTCTTTTGAAGA  
GGTATTGAACCATGATGCAAGAAAGGAGTGAGAAAAACGAGTTTCTTCTGGGGGCTCCAGATTATTATCGTCTCTTAACCTGGT  
GGGAAAAAGGCTTGAAGTGGGTATGGGTGCTTACTCTGCTCCCTATTTATCGGATTGGTTTGTCTTGTACAGTTT  
AGACGTTTATGGCTATCTTTACCTCCCTTTGCCAATACTTGGTTTTCTAGGGTTAGTTTGTCTTTCTATTATTGGAAGCTT  
CGACTAGATAATCGTGATGGTGTCTAAATGAAGCGGGAGCTGAGGTCTACTATCTCTGGACCAGTTTGAAGAAATATGTTGC  
GTGAGATTGCACGATTGGATCAGGCTGAACTGGAAAGTATTGTGGTCTGGAATCGCCTCTTGGTCTATGCGACCTTATTTGGC  
TATGCGGACAAGGTTAGTCAATTTGATGAAGGTTCTCATGATTCAAGATGGAAAAATCCAGATATCAATCTCTATGTAGCTTATGG  
CTGGCACAGTACGTTTTATCATTCAACAGCACAAATGAGCCATTATGCTAGTGTGCAAAATACAGCAAGCACCTACTCTGTAT  
CTTCTGGAAGTGAAGTTCTGGTGGTGGCTTCTCTGGAGGCGGAGGTGGCGGCAGTATCGGTGCCTTTTAA

## 4191.2 (SEQ. ID. NO. 338)

ATGAAAAAAGTAAAGATAATTTTCAAGAGGAGTTCAGGACTGTGCTGTATATCTCAGTTGACAGCTTTTTCTTCCGATAG  
TTGCTTTAGCAGAAACGCCTGAAACCAAGTCCAGCGATAGGAAAAAGTAGTGATTAAGGAGACAGGGCAAGGAGGAGCGCTTC  
TAGGAGATGCCCTTTTGAAGTTGAAAAACAATACGGATGGCACAACCTTTTCGCAAGGACAGAGGCGCAACAGGAGAAAG  
CGATATTTTCAAACATAAAACCTGGGACATACACCTTGACAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACA  
ATGGACTGTTGAAGTTGAGAAGAATGGTCCGACGACTGTCCAAAGTGAACAGGTAGAAAAATCGAGAAGAGGCTCTATCTGA  
CCAGTATCCACAAACAGGGACTTATCCAGATGTTCAACACCTTATCAGATTATTAAGGTAGATGGTTCCGAAAAAAGCGGA  
CAGCACAAGGGCGTTGAATCCGAATCCATATGAACGTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATA  
ATTTGGATGATAACCAATATGGAATCGAATTGACGGTTAGTGGGAAAAACAGTGTATGAACAAAAAGATAAGTCTGTGCCGCT  
GGATGTCGTTATCTTGGTCGATAACTCAAATAGTATGAGTAACATTCGAAACAAGAATGCTCGACGTGCGGAAAGAGCTGGT  
GAGGCGACAGCTTCTCTTATTTGATAAAATTACATCTCATGATTCAAGAAATAGGGTAGCGCTTGTGACTTATGCTCACTCTT  
TGATGGGACCGAGTTTACAGTAGAAAAAGGGGTAGCAGATAAAAAACGGAAAGCGATTGAATGATTCTCTTTTTTGGAAATTAT  
GATCAGACGAGTTTACAACCAATACCAAGATTATAGTTATTTAAAGCTGACTAATGATAAGAATGACATTGTAGAATTAA  
AAAATAAGGTACCTACCGAGGCGAGAAGACCATGATGGAATAGATTGATGTACCAATTCGGTGCCACTTTTACTCAGAAAGC  
TTTGATGAAGGCAGATGAGATTTTGACACAACAAGCGAGACAAAAATAGTCAAAAAGTCATTTTCCATATTACGGATGGTGTC  
CCAATATGTCGATCCGATTAATTTTAAATCATGCTACGTTTGTCTCATCATATAAAATCAACTAAATGCATTTTTTATGATAA  
TCTCCTAATAAAGATGGAATACTATTAAGTGAATTTTATTACGCAAGCAACTAGTGGAGAACATACAATTGTACGGCGGAGATG  
GGCAAGATTACAGATGTTTACAGATAAGACAGTTTATGAAAAAGGTGCTCTGACGCTTCCGAGCTTTCCAGTAAAAATA  
TTCTGAAATGAAGGCGGCTGGTTATGCAGTTATAGGCGATCCAATTAATGGTGGATATATTGGCTTAATTGGAGAGAGAGT  
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TAGGTGGTCCACAAAATGATGGTGGTTTGTAAAAAATGCAAAAGTCTCTATGATACGACTGAGAAAAGGATTCGTGAAC  
AGGTCTGTACCTTGAACGGATGAAAAAGTTACGTTGACCTACAATGTTTCGTTTGAATGATGAGTTTGAAGCAATAAAATTTT  
ATGATACCAATGGTGAACAACCTTACATCCTAAGGAAGTAGAACAGAACACAGTGCAGGACTTCCCGATTCTTAAGATTCTG  
TGATGTGCGGAAGTATCCAGAAATCACAATTTCAAAGAGAAAAAAGTGGTGACATTGAGTTTATTAAGGTCAATAAAAAAT  
GATAAAAAACCACTGAGAGGTTGCGGTCTTATGCTTCTCAAAAACAACATCCGGATTATCCAGATATTTATGGAGCTATTGATC  
AAAATGGCACTTATCAAAATGTGAGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGAGATGGGAAATATCGATT  
ATTTGAAAAATCTGAACCGCTGTTTATAAACCCGTTCAAAAATAAGCCTATCGTTGCTTCCAAATAGTAAATGGAGAAGTC  
AGAGATGTGACTTCAATCGTTCCACAAGATATACAGCGGGTTACGAGTTTACGAATGATAAGCACTATATTACCAATGAAC  
CTATTCTCCTCAAAGAGAGAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTGTCATGATGATGGGA  
GGAGTTCTATTATACACACGGAACATCCGTAA

## 4191.3 (SEQ. ID. NO. 339)

ATGAAATCAATCAACAAATTTTAAACAATGCTTGTGCTTATTAAGTACAGCGAGTAGCCTGTTTTTCACTGCAACAGTTTTT  
TGCGGCTGGGACGACACAACATCTGTTACCGTTCAAACTATTGGCAACAGATGGGGATATGGATAAAATTTGCAAAATGAG  
TTAGAAACAGGTAACATATGCTGGTAATAAAGTGGGTGTTTCTACCTGCAAAATGCAAAAGAAATTTGCCGGTGTATGTTCTGTT  
GGCAAAATACTAATAATGAAATTTATGATGAAAAAGGCAAACTTACAGGAGTGAATTTGATCCACAAACATTTAAACTCTC  
AGGGCAATCCCGGCAATGCAATGAAAAAATAACGAAAGTGAAGAGCTAAATTTAACACGCAAAATTTACAGCTGC  
TAAGTATAAAATTTATGAAATTCACAGTTTATCAACTTATGTGGTGAAGATGGAGCAACCTTAAACAGGTTCTAAAGCAGTTT  
CAATTGAAATTTGAATTACCATTTGAACGATGTTGTGGATGCGCATGTGTATCCAAAAATACAGAAGCAAGCCAAAAATTTGA



TAAAGATTTCAAAGGTAAAGCAAATCCAGATACACCACGTGTAGATAAAGATACACCTGTGAACCACCAAGTTGGAGATGTT  
GTAGAGTACGAAATTTGTACAAAAATTCAGCACTTGCTAATTATGCAACAGCAAACCTGGAGCGATAGAATGACTGAAGGTT  
TGGCATTCAACAAAGGTACAGTGAAGTAAGTGTGATGATGTTGCACTTGAAGCAGGTGATTATGCTCTAACAGAAGTAGC  
AACTGGTTTTGATTTGAAATTAACAGATGCTGGTTTAGCTAAAGTGAATGACCAAAACGCTGAAAAAAGCTGTGAAAACTACT  
TATTCGGCAACATTGATGACAAAGCAATTTAGAAGTACCAGAATCTAATGATGTAACATTTAACTATGGTAATAATCCAG  
ATCACGGGAATCTCCAAAGCCGAATAAGCCAAATGAAAACGGCGATTTGACATTGACCAAGACATGGGTTGATGCTACAG  
GTGCACCAATTCGGGCTGGAGCTGAAGCAACGTTGCTGATTGGTTAATGCTCAGACTGGTAAAGTTGTACAACTGTAACCTTG  
ACAACAGACAAAAATACAGTTACTGTTAACGGATTGGATAAAAAATACAGAATATAAATTCGTTGAACGTAGTATAAAAGGG  
TATTCAGCAGATTATCAAGAAATCACTACAGCTGGAGAAATGCTGTCAAGAACTGGAAAGACGAAAAATCCAAACCACTT  
GATCCAACAGAGCCAAAAAGTTGTTACATATGGTAAAAAGTTTGTCAAAGTTAATGATAAAGATAATCGTTTAGCTGGGGCAG  
AATTTGTAAATTGCAAATGCTGATAATGCTGGTCAATATTTAGCACGTAAAGCAGATAAAGTGAGTCAAGAAGAGAAGCAGTT  
GGTTGTTACAAACAAAGGATGCTTTAGATAGAGCAGTTGCTGCTTATAACGCTCTTACTGCACAACAAACAACTCAGCAAGAA  
AAAGAGAAAGTTGACAAAGCTCAAGCTGCTTATAATGCTGCTGTGATTGCTGCCAACAATGCATTTGAATGGGTGGCAGATA  
AGGACAATGAAAAATGTTGTGAAATTAGTTTCTGATGCACAAGGTGCGTTTGAATACAGGCCTTCTTGCAGGTACATATTAC  
TTAGAAGAAACAAACAGCCTGCTGGTTATGCATTACTAAGTACGCGTCAAGAAATTTGAAGTCACTGCAACTTCTTATTCAGC  
GACTGGACAAGGCATTAGATATACTGCTGGTTAGGTAAGATGACGCTACAAAAGTAGTCAACAAAAAATCACTATCCC  
ACAAACGGGTGGTATTGGTACAATTATCTTTGCTGTAGCGGGGCTGCGATTATGGGTATTGCAGTGATCGCATATGTTAAA  
AACAACAAAGATGAGGATCAACTTGCTTAA

## 4191.4 (SEQ. ID. NO. 340)

ATGACAATGCAGAAAATGCAGAAAATGATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGTATGGGGTGCACA  
TGCAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAAGTATCAGGAGGTGGTTAGTCAATTGCCATCTCGT  
GATGGTATCGGTTGCAAGTATGGAAGTTGGATGATTGCTATTCCTATGATGATCGGGTGCAAATGTAAGAGACTTGCATTC  
GTGGGATGAGAAATAAACTTCTTCTTTCAAAAAGACTTCTGTTGATGACCTTCTTGAGAATCAGATTGAAGTATCTCAT  
TTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTTATCCAGCTGAATTTCTTTTGAATGACAG  
ATCAAAACGGTAGAGCTTTGGTCAATTGTAGCGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATC  
AAGAACCAATCGCTTGGAGGGTGTGCGCTTAAATTTGGTATCAGTAGCAAGAGATGTTTCTGAAAAAGAGGTTCCCTTGAT  
TGGAGAATACCGTTACAGTTCTTCTGGTCAAGTAGGGAGAAGTCTCTATACTGATAAAAAATGGAGAGATTTTGTGACAAAT  
CTTCTCTTGGGAAGTATCGTTTCAAGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGCT  
GGTAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTGACTTTATGAAGGTGGATGGT  
CGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAAGTCAAGTGAAGAAGAAAGCGGACACTATACTCTGTTCTTCAAAATG  
GTAAGGAAGTAGTTGTAACATCAGGGAAGATGGTCTGTTCCGAGTGAAGGTCTAGAGTATGGGACATACTATTTATGGGA  
GCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTCTTTACAATCGGGAAAGATACCGTAAGGAAGTGGTAA  
CAGTGGTTAAAAATAACAAGCACCAGGATTGATGTCCAGATACAGGGGAAGAAACCCCTGTATATCTTGATGCTTGTG  
CCATTTTGTGTTTGGTAG

## 4191.5 (SEQ. ID. NO. 341)

ATGAGCCACATATACTTATCTATTTTCAAGTCTCTTGTGATGCTAGGACTTGTCAATGTTGCTCAAGCCGATGAATATTTA  
CGCATCGGTATGGAAGCAGCATATGCTCCCTTAACTGGACCCAGGATGATGATAGCAACGGAGCTGTCAAAATCGATGGGA  
CCAATCAGTATGCCAACGGATACGATGTTCAAAATCGCAAGAAAAATCGTAAGGACTTAGGTAAGAAGACCTTTGGTTGTAA  
AACCAAGTGGGAAGGTCTAGTCCCTGCCCTTACTTCTGTAAGATTGACATGATTATCGCAGGTATGAGTCCAACTGAGAA  
CGCAACAAGAAAATGCTTTTTCGAGCAGTTACTATACTAGCGAACCAAGTTTGTCTGTCAAAAAAGATTCTGCCTACGCAA  
GTGCTAAATCTTTGGATGACTTTAACGGTGCAAAAAATCACTTCTCAACAAGGGGTCTACCTTTATAACTTGATTGCACAAATC  
CCAGGTGCTAAAAAGAAACAGCCATGGGAGACTTTCGCTCAATGCGACAAGCTCTTGAGGCTGGTGTCAATTGATGCTTATG  
TTTCTGAACGTCCAGAACGACTGACTGCTGAAGCTGCGAACTCAAGTTCAAGATGATTCAAGTAGAACCTGGTTTCAAAAC  
TGGGGAAGAAAGATACAGCTATCGCTATCGGGCTTCGTA AAAATGACAATCGTATTAGCCAAATCAATGCCAGCATTGAAACC  
ATTTCAAAAAGATGACCAAGTTGCCCTTGATGGATCGTATGATCAAGGAACAACCTGCCGAAGCTACAACAAGTGAAGAGACTA  
GCAGTAGTTTCTTTAGCCAAGTTGCTAAAAATCTTTTGTGAAAACCTGGCAACAACCTTTGCGTGGTGTGATCATCTTTTAA  
TCTCTATCGTCGGAACCATCATAGGTCTCATTATTGGACTTGCCATTGGTGTCTTCCGTAAGTCTCTCTCTGAAAACAAAG  
TCATTTACGGCCTACAAAACTAGTCGGCTGGGTTCTCAATGCTACATTGAAATTTCCGTGGTACGCCAATGATTGTTCAA  
TCGATGGTTATCTACTATGGAACCTGCCAAGCTTTCCGGATCAACC  
TTGACCGTACACTGGCTGCTATCTTCACTGTTTCAATAACCCGTTGCCCTACATGACTGAAATCGTCCGTGGTGGTATCCTA  
GCAGTTGACAAGGGACAATTTGAAGCTGCGACTGCTCTTGGTATGACCCATAACCAGACCATGCGTAAGATTGTCCTACCTC  
AGGTAGTCCGTAACATCCTACCTGCAACTGGTAATGAATTTGTCAATATCAAGATACATCTGTATTGAACGTTATCTCT  
GTTGTGCAACTTTATTTCTCAGGAAATACCGTGGCAACACAAACCTATCAATACTCCAGACATTTACAATCATCGCCGTGAT  
TTACTTTGCTCTACCTTACCGTAACACGATCTCTACGCTTTATCGAGCGCAGAATGGACATGGATACCTACACTACAGGTG  
CTAACCAATGCAACGGAGGATTGAAATAA

## 4191.6 (SEQ. ID. NO. 342)

ATGACACAAGCAATCCTTGAAATTAACACCTCAAAAAATCCTATGGACAAAACGAAGTGCTAAAAGACATTTCACTCACTG  
TCCACAAGGGAGAGGTATCTCTATCATCGGAAGCTCTGGAAGCGGAAATCGACCTTCTACGCTCCATTAACCTACTTGA  
AACACCAACTGATGGACAAATCCTTTATCATGGACAAAACGCTCTGAAAAAGGCTATGACCTCAGCAATACCGTGAAAAAG  
TTGGGGATGGTTTTTCAATCCTTTAACTCTTTGAAAAATCTCAATGTTCTTGAAAAACAAATCGTCTGCTCAGACAACTGCTC  
AAAACGCGAACGACAGAAAGCTGAAAAGATTGCCAAAGAAAACCTGGAAAAGGTGCGCATGGGAGAACGCTACTGGCAAG  
CCAAACCAAAAAACACTCTCAGGTGGTCAAAAAACAGTGTGGCCATCGCTCGTCCCTCTCCATGAATCCGGACGCTATTTCT  
CTTTGATGAACCAACATCAGCTCTCGATCCAGAAATGGTTGGGAAGTCTCAAAATCATGCGAGGATGCTCAGGAAGGC  
TTGACCATGATTGTCGTAACCCATGAAATGGAATTTGCCCGTGATGTCTCTACCGTGTTATCTTTATGGATAAGGGCGTGAT  
CGCTGAAGAAGGTAAACCAGAAGACCTTTCACCAATCCTAAAGAAGACCGAACAAGAGTTCTTCAACGCTATCTCAA  
ATAA

## 4192.3 (SEQ. ID. NO. 343)

ATGAAAAAGTATCAACTTCTATTCAAAATAAGTGCAGTCTTCTCTTACTTATTTTTTCGTAATTTAGTCTTTCTCAGCTGACGCTT  
ATCGTCCAAACTATTGGCAATTTTCTCTCAGATAGGCAATTTATCTGGATTCAAAATATCTTGAGTTTACTTTTTATTGGA  
GTCATGATTGTGGTTCTTGTAAAGACAGGCCATGGTTATCTCTTCCGCTTCCAAAGAAAAAATGGCTTTGGTATTCGATTTT  
GACAGTATTAGTGCTAGTGTTCCAGATCTTTTTAACGTTTCAGACAGCTAAACATGTTTCAGTCAACTCGCGAAGGTTGGGCTG

TATTGATTGGTTATAGTGGGACTAACTTTGCAGAGCTAGGTATTTATATAGCCCTGTTCTTTCTGGTTCCACTGATGGAAGAA  
TTGATTATAGAGGATTACTGCAACATGCTTTCTTTAAGCATTTCGCGATTTGGTCTTGATTGGCTTCTTCTTCTATTTTATTTG  
CTCTCCCTCATTTTCAAGCCTGCCTAGTCTGTTAGATATCTTCGTCTTTGCAACAGTTGGAATCATCTTTGCTGGTTTGACCC  
GCTATACCAAGAGCATTTATCCATCTATGCGGTGCATGTGATCAATAATATTGTAGCGACCTTCCCGTTTTTGCTCACTTTTC  
TACATAGGGTCTTGGGGTAA

## 4193.1 (SEQ. ID. NO. 344)

ATGAACAAGAAACAATGGCTAGGTCTTGGCCTAGTTGCAGTGGCAGCAGTTGGACTTGGCTGCATGTGGTAACCGCTCTTCTC  
GTAACGCAGCTTCATCTTCTGATGTGAAGACAAAAGCAGCAATCGTCACTGATACTGGTGGTGTGATGACAAATCATTCAA  
CCAATCAGCTTGGGAAGGTTTGCAGGCTTGGGGTAAAGAACACAATCTTTCAAAAGATAACGGTTTCACTTACTTCCAATCA  
ACAAGTGAAGCTGACTACGCTAACAACTTGCAACAAGCGGCTGGAAGTTACAACCTAATCTTCGGTGTGGTTTGGCCCTTA  
ATAATGTCAGTTAAAGATGCAGCAAAAGAACACACTGACTTGAACCTATGTCTTGATTGATGATGTGATTAAAGACCAAAAGAA  
TGTTGCGAGCGTAACTTTTCCTGATGAATGAGTCAGGTACcTTGCAAGTGTGGCTGCAGCAAAAACAACTAAGACAAAACAA  
GTTGGTTTTGTAGGTGGTATCGAATCTGAAGTTATCTCTCGTTTTGAAGCAGGATTCAAGGCTGGTGTGCGTCAGTAGACCC  
ATCTATCAAAGTCCAAGTTGACTACGCTGGTTTCATTTGGTGTATGCGGCTAAAGGTAAAACAATTGCAAGCCGCACAATACGCA  
GCGGTGCAGATATTGTTTACCAAGTAGCTGGTGGTACAGGTGCAGGTGTCTTTGCAGAGGCAAAATCTCTCAACGAAAGCC  
GTCCTGAAAATGAAAAAGTTTGGGTTATCGGTGTGATCGTGACCAAGAAGCAGAAGGTAATACTTCTAAAGATGGCAA  
AGAATCAAACCTTTGTTCTTGATCTACTTTGAAACAAGTTGGTACAACCTGTAAGATATTCTAACAAGGCAGAAAGAGGA  
GAATTCCTGGCGGTCAAGTGATCGTTTACTCATTGAAGGATAAAGGGGTTGACTTGGCAGTAACAAACCTTTCAGAAGAAG  
GTAAAAAGCTGTGCAAGATGCAAAAGCTAAAAATCCTTGATGGAAGCGTAAAAAGT  
TCCTGAAAAATAA

## 4193.3 (SEQ. ID. NO. 345)

ATGTCTAAAAAATTACAACAAATTCGGTTCCTTGATTTCTGTATTCTAGGAATTTTACTCGGAGCCATTGTCATGTGGATC  
TTCGGTTATGATGCTATTTGGGGCTACGAAGAATTGTTCTATACAGCCTTTGGCAGTCTGCGTGGGATTGGAGAAAATCTTCCG  
TGCTATGGGTCCTCTGGTCTTGATTGGTCTTGGTTTTGCCCTTGCCAGTCGAGCTGGTTTTCTTAAAGCTCGGACTTCTGGTCA  
GGCTTTGGCAGGTTGGATTCTCAGTGGTTGGTTTTGCCCTGTCGATCCAGATATGCCCGTCCCTTGATGATTCTAGCAACCA  
TCGTGATTGCCTTGATTGCTGGTGGGATTGTCGGAGCGATTCCAGGTATTCTTAGGGCCTATCTAGGGACGTCAGAGGTTATT  
GTAACCATCATGATGAACCTACATTGTCTTGATGTAGGGAATGCCTTTATCCATGCTTCCCTAAAGACTTCATGCAAGTAC  
AGATTGACCATTCGTGTGGGCTAATGCAACCTATCAGACACCTTGGTGGCTGAGTTGACTGGTAACCTACGGATGAAT  
ATTGGTATTTTCTTTGCCATCATTGCCGTTGCAGTTATTTGGTTTCATGCTCAAGAAAACAACCTTGGTTTTGAAATCCGTGCA  
GTTGGTCTTAATCCACATGCTTCAGAATATGCTGGTATTTCTGCCAAGCGGACTATTATCTATCTATGATTATTTACGGTGCC  
TTGGCAGGTCTTGGTGGAGCTGTTGAAGGTTTGGGAACCTTCCAGAACGCTATGTTCAAGGTTTCGTCAATTAGCTATCGGATT  
TAACGGAATGGCGGTTAGTTTGGCTTGGGCCAACTACCAATTTGGTATACCTTTTGCAGCCTTCTATTGGCGTTCTCCAAGT  
TGGGGCTCTGGTATGAATGCGGCGCAGGTACCATCTGAGCTTGTGAGCATTGTAACAGCGTCTATTATCTTCTTTGTCAGTG  
TTCATTACCTTATCGAACGCTTTGTCAAACCGAAAAAACAAGTTAAAGGAGGTAAGTAA

## 4194.1 (SEQ. ID. NO. 346)

ATGGGAGTGAAAAAGAACTAAAGTTGACTAGTTTGGCTAGGACTGTCTCTGTTAATCATGACAGCCTGTGCGACTAATGGGG  
TAACTAGCGATATTACAGCCGAATCGGCTGATTTTGGAGTAAATTTGGTTTACTTCTTTGCGGAAATCATTCCGTTTTATCGT  
TTGATATTAGTATCGGAGTGGGGATTATTCTCTTTACGGTCTTGATTCTGACAGTCCCTCTTGCCAGTCTTTCAGGTGCAATGG  
TGGCTTCTAGGAAAAATGCAGGAAGCTCAGCCACGCATTAAGGCGCTTCGAGAACAATATCCAGGTGAGATATGGAAAGCA  
GAACCAAACCTAGAGCAGGAAATGCGTAAAGTATTTAAAGAAATGGGTGTCAGACAGTCAGACTCTCTTTGGCCGATTTTGAT  
TCAGATGCCGTTATTTTGGCCCTGTTCCAAGCCCTATCAAGAGTTGACTTTTTAAAGACAGGTCAATTTCTTATGGATTAACTT  
TGGTAGTGTGGATACAACCCTTGTCTTCCGATTTTAGCAGCATTTTCACTTTTAAAGTACTTGGTTGTCCAACAAAGCTTT  
GTCTGAGCGAAATGGCGCTACGACTGCGATGATGTATGGGATTCCAGTCTTGATTTTTATCTTTGAGTTTATGCGCCAGGTG  
GAGTCGCCCTATACTGGACAGTGTCTAATGCTTATCAAGTCTTGCAAAACCTATTTCTTGAATAATCCATTCAAGATTATCGCA  
GAGCGGAGGCCGTAGTACAGGCACAAAAAGATTGAAAAATAGAAAAAGAAAGCCAAGAAAAAGGCTCAGAAAAACGAA  
ATAA

## 4194.4 (SEQ. ID. NO. 347)

ATGGTTATCGATCCATTITGCTATCAACGAACTAGACTATTACTTAGTTTACACTTCCACAGTGATCATATCGACCCATACAC  
AGCTGCAGCAATTCTCAATAATCCTAAGTTAGAGCATGTTAAGTTTATCGGTCTTACCCTGTGGACGAATCTGGGAAGGAT  
GGGGTGTTCAAAAAGACGTATCATCGTTGTTAAACCAGGTGACACTATCGAATTAAGATATGAAGATTTCATGCAGTAGA  
ATCATTTGACCGTACTTGGTGGTAACTCTCCAGTGAACGGTGCTGATGAGACAGGCGGTGAACCTTGCTGGCTTGGCTGTTA  
CAGATGAAGAAATGGCTCAAAAGGCTGTAACTATATCTTTGAAACACCAGGTGGAACCATCTATCATGGTGCAGATTCTCA  
CTTCTCAAACTATTTTGCAAAACATGGTAAAGACTTTAAATTTGATGTTGCTTTGAATAACTATGGTGAATAATCCGGTAGGTA  
TCCAAGACAAAATGACATCTATCGACCTTCTTCGTATGGCAGAAAACTCGGTACCAAAGTCAATATCCAGTTCACTATGAT  
ATCTGGTCTAACTTCATGGCTTCTACTAATGAGATTCTAGAACTTTGGAAAAATGCGAAAAAGATCGCTTGCAATACGATTTC  
TCCATTTATCTGGGAAGTTGGCGGTAAGTACACTTATCTCAAGATCAACACTTAGTAGAATACCATCATCCACGTGGTTTTG  
ATGATTGTTTTGAACAAGACTCTAACATTCAATTTAAAGCTTTGCTATAA

## 4196.2 (SEQ. ID. NO. 348)

ATGTTCTTTTCAAGGCTGGTTGTCTAGTTTGGCTAACTTATATCCATGATTTACTGGGGGTTCTTTTCCAGATAGTCCATTTT  
TAAATGCCCTTTGAAAGTGCTATTGCGGCTCCTTTGGTAGAAGAACCTTGAAATTTGTCACCTGTTTTGTTTTGGCTTTGA  
TTCTGTGCGAAAAATAAAACTTTTGTCTTTTACTTGAATTTGCTTCCGTTTGGGATTCCAAATGATTAAGGATATTGGTTATA  
TTCTGTACGGATTGGCAGAGGCTTTGACTTTACTATTTCCGCAATTTTAGAGCGTATCATCTCAAGGAATTGCCCTCACTGG  
ACTTTTTCAGGTCTAGCTGTAGTAGGTGTTTACTTGTCTTACAGAGCCTATAAAGGACAGAAGGTTGGCAAGAAACAGGGCC  
TTATTTTTCTAGGTTTAGCCTTTGGGAACCTCACTTCTTGTTTAACTCTCCTTTTGTGGAGTTGGAAACAGAGTTGCCTTTAGCGA  
TTCCAGTGGTTACGGCTATTGCTCTCTATGGTTTTATCATGCTTATTGCTTTGTTGAGAAACACAATGAGTTGATGACCTAG

## 4197.1 (SEQ. ID. NO. 349)

ATGAAGGTGGAACACGCTTGGCAGCTCCTTTTCGAGGATGTGCGATTTTTTTTATTAGGATACTAATTATGGAGTTGCAAGAATT  
 AGTGGAGCGCAGTTGGGCAATCCGACAAGCTTATCACGAAGTGAAGTTAAGCATCATGATTCCAAGTGGACGGTGAAGA  
 AGACCTCTTGGCTTTATCTAATGATATTGAAAATTTCCAACGACTGGTGATGACAAAGCAAGGACGCTACTATGATGAAACA  
 CCTACACACTGGAACAAAAAATTTTCAGAAAAATATCTGGTGCTATTAGAACTTTCTCAACGTTTGGATATAGACATTCTGAC  
 GGAATGGAACCTTCTCTCTGATAAAGAAAAAGCAATTGAACGTTAGGACTTGGAAAGTAG

## 4197.4 (SEQ. ID. NO. 350)

ATGCTTGATTGGAAACAATTTTTCTAGCCTATCTGCGCTCCCGTAGTCGCTTTTTATCTATCTGCTTTCTTTGGCATTTCCTTG  
 TCTTACTCTTTTCAGTTTTTATTGGCAGTCTAGGAATTTACTTCCTCTACTTTTTCTTCTTGTTGGCTTTGTAACCATATTATTT  
 TTCACTTGGGACATATTGGTGGAAACGCAGGTCTATCGCCAGGAACCTTCTATGGAGAGAGGGGAAGCCAAGTCTCCTTTGG  
 AAATAGCTTTAGCAGAAAAATTAGAAGCGCGTGAGATGGAACCTCTATCAGCAGAGGTCAAAGCAGAAAGAAAACTGACGG  
 ATTTGCTGGATTACTATACCTTGTGGGTCCATCAGATAAAGACCCCCATTGACCCAGTCAACTCTTAGTTGCAGAAAGTGGTC  
 GACCGCCAACTGAAGCAGCAGCTAGAACAGGAAATTTTCAAATCGACTCTATACCAACCTAGTTTTACAGTACCTGGCTT  
 TAGAAAGTTTCCATGATGATTGGTCTTAAAGCAGGTTCAAATTGAGGACTTGGTCAAGGAAATAATTCGTAATATGCTCTT  
 TTCTTTATTCAAAGGGCTAAATGTCAATCTACATGACCTTGATAAAGAAATCGTGACGGATAAAAGTGGCTGCTAGTGG  
 TTATTGAGCAAAATCATCTCAACAGTCTCAAGTACACCAAGGAAGGTGGTCTGGAGATTATATGGATGACCAAGAGCTTTG  
 TATCAAAGATACGGGAATCGGGATAAAAAACAGTGATGTCTCCGAGTATTTGAACGTGGCTTTTCAGGATACAATGGCCGT  
 TTGACCCAGCAGTCTCTGGACTTGGCCTTATCTATCTAAGAAATTTCTGAAGAACTGGGGCACCAGATTCTGATCGAGTC  
 TGAGGTGCGAAAGGAACGACAGTGCGGATTCAAGTTGCTCAAGTGAACCTTAGTCCTTGAGTAA

## 4211.2 (SEQ. ID. NO. 351)

ATGGAACCTTAATACACACAATGCTGAAATCTTGCTCAGTGACGTAATAAGTCCCACTATCCGCAGGATGAACTGCCAGAGA  
 TTGCCCTAGCAGGGCGTTCAAATGTTGGTAAATCCAGCTTTATCAACACTATGTTGAACCGTAAGAATCTCGCCCGTACATCA  
 GGAACAACTGGTAAAAACCCAGTCTCTGAACTTTTTAACTATTGATGACAAGATGCGCTTTGTGGATGTGCTGGTTATGGCTA  
 TGCTCGTGTCTTAAAAAGGAACGTGAAAAAGTGGGGGTGCATGATTGAGGAGTACTTAACGACTCGGGAAAAATCTCCGTGCG  
 GTTGTCAGTCTAGTTGACCTTCGTCATGACCCGTCAGCAGATGATGTGCAGATGTACGAATTTCTCAAGTATTATGAGATTCC  
 AGTCATCATTTGTGGCGAAGGCGGACAAGATTCTCTGGTGAATGGAACAAGCATGAATCAGCAATCAAAAAGAAATTT  
 AAACCTTTGACCCGAGTGACGATTTTCATCTCTTTTCATCTGTCAAGGACGGGATGGATGAGGCTTGGGATGCAATCTTAG  
 AAAAATTGTGA

## 4211.3 (SEQ. ID. NO. 352)

ATGACAAAGAAACAACCTTCACTTGGTGATTGTGACAGGGATGAGTGGCGCAGGGAAAACTGTAGCCATTCAAGTCCTTCGAGG  
 ATCTAGGTTATTTACCAATTGATAATATGCCGCCAGTCTCTTGCCTAAGTTTTTGACGCTGGTTGAAATTAAGGAAGACAAT  
 CCTAAGTTGGCCTTGGTAGTGATATGCGTAGCCGTTCTTTCTTTTCAGAGATTCAAGCTGTTTTGGATGAGTTGGAAAAATCA  
 AGATGGTTTTGGATTTCAAATCCTCTTTTGGATGCGGCTGATAAGGAATTGGTGGCTCGTTACAAGGAAACCAGACGGAGT  
 CACCCACTAGCAGCAGACGGTCGTATTTAGATGGAATCAAGTTGGAACGTGAACCTTTGGCACCTTTGAAAAATATGAGCC  
 AAAATGTGGTGGATACGACTGAACCTCACTCCACGTGAGTGCGCAAAACCCCTTGACAGCAGTTTTCAGACCAAGAACAAGC  
 CCAGTCTTTCCGTATCGAAGTCATGTCTTCGGATTAAAGTATGGAATCCCGATTGATGCGGACTTGGCTTTGATGTCCGTTT  
 CTTGCCAAATCCCTATTATTTACCAGAACTGAGAAACCAACGGGTGTGGATGAACCTGTTTATGATTATGTATGAACCATC  
 CTGAGTCAGAAGACTTTTATCAACATTTATTTGGCCTTGATTGAGCCGATTCTGCCAAGTTACCAAAAGGAAGGTAAGTCCGTT  
 TTGACCAATTGCCATTGGATGTACGGGTGGACAACCCGATTCTGGCATTTGCTAAACGCTTGGCGCAGGACTTATCCAAGA  
 ATTGGTCTGTTAATGAAGGGCATCGGCACAAAGACCCGACAGAAAGGAACCGTAAACCGTTTCATGA

## 4211.4 (SEQ. ID. NO. 353)

ATGAGAAAACCAAAGATAACGGTGATTGGTGGAGGGACTGGAAGTCCCGTCATTCTAAAAAGTCTGCGGGAAAAAGATGTG  
 GAAATCGCAGCTATCGTGACGGTGGCAGATGATGGTGGTTCTTCAGGTGAACCTCGAAAAAATATGCAACAGTTGACACCGC  
 CAGGTGATCTTCGTAATGTCTTGTGGCCATGTCCGATATGCTTAAGTTTTATGAGAAGGTCTTTTCAGTATCGGTTCTCTGAG  
 GATGCCGAGCCCTTGGTGGCCATCCATTGGGAAATCTCATTTTTCGCTTGGCTTGTGAGAAATGCAAGGTTCAACCTATAATGC  
 CATGCAGTTATTGAGCAAAATTTTCCATACAACAGGGAAAAATTTATCCTTCCAGTGACCATCCTTTGACCCCTTCATGCAGTCTT  
 TCAGGATGGGACAGAAAGTGGCTGGAGAGATCATATTGTAGACCATCGAGGCATAATTGACAATGTCTATGTGACCAATGCC  
 CTAACGATGATACGCCCTCTGGCCAGCCGTCGAGTAGTGACACCATCCTTGAAAGTGACATGATTGTCTAGGGCCAGGTT  
 CCCTCTTTACCTCTATTTTGGCCAAATATCGTGATTAAAGGAAATTTGGCGGGCTCTTTTGGAAACCAAGGCAGAAATTCCTAT  
 GTCTGCAATATCATGACCCAACGTGGGGAGACGGAACACTTTACAGATAGCGACCACGTGGAAGTCTTGTCATCGTCACCTTG  
 GTCGCCCTTTTATCGACACTGTCTTGGTGAATATTGAAAAAGTGCTCAGGAATACATGAATTTCAACCGTTTGTGATGAATAC  
 TTAGTGCAAGTGGAAACAGATTGTAGGTCTTTGTAAGCAAGTTTCGCGCGTGATTTCATCTAACTTCCTTCGTCTGGAAAA  
 TGGCGGTGCCTTCCACGATGGAGATTTGATTGTGGACGAGTTGATGCGCATTATACAGGTGAAAAAATGA

## 4213.1 (SEQ. ID. NO. 354)

ATGAAAAATTTGATAAAGTTGCTAATAATTAGATTGATTGTTAACTTAGCAGACAGTGTATTTTATATAGTAGCATTGTGGCA  
 CGTTAGCAATAATTATCTTCGAGCATGTTCTTAGGAATATTTATTGCAGTAAATTTATCTACCGGATTGTTACTAATCTTTTT  
 TGGACCAAGTTATTGACAGAGTAAATCCGCAAAAAATCTTATAATATCAATTTTGGTTCAATTAGCAGTGGCTGTAATATTTT  
 TATTATTATTAACCAAATATCATTTTGGGTGATAATGAGTCTAGTGTTTATTTCAGTAATGGCTAGCTCCATAAGTTACGTG  
 ATAGAAGATGTGTTGATTCTCAAGTGGTAGAATATGATAAGATTGTATTGCAAAATCTCTTTTTAGTATTTTCGTATAAAGT  
 ATTAGATTCTATTTTAAATTCATTCCGATCATTTTACAGGTGGCAGTAGGATTTATTTTATTGGTTAAGATAGATATAGGCAT  
 ATTTTACTTGCTCTATTTATATTGTTGTTGTTAAAAATTTAGAAGTGAATGCGAATATAGAAAACTTCTCTTTCAAATATTA  
 CAAGAGAGAAGTGTGCAAGGTACAAAGTTATTTTAAATAATAAATTTATTTTAAACCAAGTATTCTTTTAAACGCTTATAA  
 ACTTTTTTTATTCATTTTCAGACAGTAGTTGTACCGATTTTTTCTATTTCGATATTTTGTGTTCCGATTTTTTATGGTATTTTTTTA  
 ACTATTGCTGGTTTGGGTGGTATATTGGGAAATATGCTAGCGCCAATCGTAATAAATTTTAAAAATCGAATCAAAATGTTGGT  
 TGTATTCTTTTTTTGAACGGCTCAAGTTGGTTAGTGAATGTTATAAAGACTATACTTTATCACTATTTTATTTTTCGTT  
 TGTTTTATGTCTAAAGGAGCTTCAATATTATTTTAAATCGTTGACCAAAATACCTCCACATCAACTCTTGGTAGGGTA  
 AATACTACCATTGATTCTATTATTTCTTTTGGAAATGCCAATTGGTAGTTAGTTGCAGGAACGCTTATTGATTGAATATTGAA  
 TTAGTGTTAATTGCTATTAGCATACCTTATTTTTTGTCTTATATTTTTTATACGGATAATGGATTGAAAGAATTTAGTATAT  
 ATTAG

## 4213.2 (SEQ. ID. NO. 355)

ATGATGTCTAACAAAAATAAGGAAATCTGATTTTTGCGATTCTCTATACAGTCCTCTTTATGTTTGATGGCGTTAAATTTGCTG  
GCTTCTTTAATGCCATCTGCCATTGCAAAATATCTTGTATGTAGTTTTAGCTCTATATGGCTCCTTCTTGTTCAGGATAGAT  
TGATCCAACAAATGGAAGGAGATTAGAAAGACTAAAGAAAAATCTTCTTTGGAGTCTTAACAGGATGGCTCTTTCTCATTCT  
GATGACTGTTGTCTTTGAATTTGTATCAGAGATGTTGAAGCAGTTTGTGGGACTAGATGGACAAGGTCTAAATCAGTCTAATA  
TTCAAAGTACCTTTCAAGAACAACCACTACTGATAGCTGTTTTGCTTGTGTCATTGGACCTCTGGTAGAAGAATTATTTTCC  
GTCAGGTCTTATTGCATTACTTGCAGGAACGGTTGTCAGGTTTACTAAGCATTATTCTGGTAGGACTTGTTTTTGCTCTGACTC  
ATATGCACAGTTTGGCTCTATCAGAGTGGATTGGTGCACTTGGTTACTTAGGTGGAGGCCTTGCCTTTTCTATTATTTATGTGA  
AAGAAAAAGAGAATATCTACTATCCCTACTTGTTCACATGTTAAGCAACAGCCTCTCCTTAATCATTTTAGCTATCAGTATA  
GTAAATGA

## 4224.1 (SEQ. ID. NO. 356)

TTGAAAAAGCCAATTATCGAATTCAAAAACGTCTCTAAAGTTTTTGAAGACAGCAACACCAAGGTTCTCAAAGACATCAACT  
TTGAGTTGGAAGAAGGGAAATTTCTACACCTTCTAGGTGCATCTGGTTCGGGGAAATCAACTATCCTAAACATTATTGCAGG  
TTTACTGGATGCGACGACAGGAGATATCATGCTAGACGGTGTTCGTATCAATGATATTCCAACCAACAAGCGCGACGTACAT  
ACCGTCTTCCAATCCTATGCCTTGTTCACACATATGAATGTGTTGAAAATGTTGCCTTTCCACTTCGCTTGCCTAAAAATTGAT  
AAGAAAGAAATCGAGCAGCGTGTAGCGGAAGTTCTCAAGATGGTTCAGTTGGAAGGTTATGAAAAACGTTCCATCCGCAAA  
CTTTCTGGAGGACAACGTCAGCGTGTGCCATCGCCCGTGTATCATCAACCAACCCCGTGTGGTCTTGTGGACGAGCCTTT  
ATCAGCGCTGGACTTGAAATTTGAGAACAGACATGCAGTACGAATTGCGTGAATTACAACAACGATTGGGCATTACCTTTGTC  
TTTTGCTACTCAGATCAGGAAGAGCTCTTGCCATGAGTGACTGGATTTTCGTTATGAATGATGGCGAGATTGTCCAGTCTGG  
AACCCTGTGGACATCTACGATGAGCCAATCAACCACCTTGTGGCCACCTTTATCGGGGAGTCAAACATCTTGCCAGGTACCA  
TGATTGAGGACTACTTGGTCTGAATTTAACGGCAACCGCTTTGAAGCGGTTGATGGTGGGATGAAGCCAAATGAACCTGTTGA  
GGTCTGTTATTCGTCCAGAGGACTTGCCTATTACCTTCTGAAGAAGGCAAGCTCCAAGTTAAGGTGATACCCAGCTTTTCC  
GTGGAGTTTCAATGAATTTATCGCTATGACGAACCTTGAAATGAATGGATGATCCACTCAACCCTAAGGCTATCGTGGG  
TGAGGAAATCGGTCTGGACTTTGAACCAGAAAGACATCCACATCATGCTCTCAATGAACCCGAAGAAGAGTTCGATGCTCGT  
ATTGAGGAGTACGTAGAAATCGAAGAGCAAGAAGCAGGTTTGATCAATGCAATCGAGGAGGAAAGAGATGAAGAAAACAA  
GCTCTAA

## 4252.1 (SEQ. ID. NO. 357)

ATGAAATCAATGAGAATCTTATTTTTGTTAGCTTTAATTCAAATCAGTTTGAGTAGCTGTTTCCTATGGAAGGAATGCATCTT  
GTCCTTTAAACAAAGTACAGCTTTTTTCATCGGAAGCATGGTTTTCGTTTCAGGAATCTGTGCTGGAGTAAATTTATCTTTATAC  
CCGTAAGCAAGAAAGTCCATAGTGTCTAGCCAGTAAGAAGTCGGTGAAGCTTTTTTACAGTATGTTACTCTTAATTAATTTGT  
TAGGAGCTGTTCTGTTTTGTCAGATAACTTGTTCATCAAAAATACGCTGCAGCAAGAATTAGTTGACTTTTTATTGCCATCCT  
TCTTTTTCTATTGTTGGCTAGATTTGCTGATTTTTTACCCTTGAAAAAATACGTGCGCGATTTTCTTGCTATGCTGGACAGAA  
AAAAGACAGTGTGGTGACTATTTAGCAACACTTCTTTCTTAAGAAATCCAATGACCATTGTCTCACTTCTGATTTATATTG  
GACTGGGCTGTTTTTTCAGCCTATCTTGTCCCAAATTCGGTTAAGAAGGAAGTTTCCTTTATGGTCAATTTTCCGAGATC  
TTGATTGGTCATTGTTACGCTCATTTTCTTTTAG

## 4252.2 (SEQ. ID. NO. 358)

ATGGTTAAAAAAATTAATTGGAATGGTGCTAGCTTTACTTTCTGTAAGTGTAGTAGGAGTAGGTGTTTTTGTCTTATACTATTTAT  
CAACAAGGGACAGAAACCTTAGCTAAAACCTATAAAAAAATCGGTGAAGAAACCAAGGTTATTGAAGCGACTGAACCTCTA  
ACCATCTGTTAATTGGGAGTGGACACCGGAAATGTTGAACGAACTGAAACTTGGGTCGGTAGAAGTGATAGCATGATCTTGA  
TGACAGTGAATCCTAAAACGAAAAAACAACAAATGATGAGTTTAGAGCGGGATATTCTGACGCGCATTTGAATCAGGGAATG  
GTCAGGCTCATGAAGCGAAACTGAACTCAGCATATGCAGATGGTGGAGCAGAGCTTGCTATAGAAACCATTCAAAAATGA  
TGAATATCCATATTGATCGCTATGTGATGGTCAATATGAGAGGATTGCAAAAACCTAGTGGATGCAGTAGGAGGTATTACAGT  
CAATAATATCCTAGGTTTTCCCAATTTCTATCAGTGACCAAGAAGAATTTAATACTATTTCTATCGGTGTTGGGGAGCAACATA  
TTGGGGGAGAAGAAGCCCTAGTCTATGCACGAATGCGTTACCAAGATCCTGAGGGGGATTATGGTCTGCAAAAACGTCAAC  
GTGAAGTTATTCAAAAAGTCATGGAAAAAGCTCTCAGTTTAAATAGCATTGGTCATTATCAAGAGATTCTAAAAGCTTTGAG  
TGACAATATGCAGACCAATATTGATTTGTCTGCAAAAAGTATCCCTAAGTTGCTAGGCTATAAAGATTCAATTTAAACCATTTG  
AAACTGCAGGTGACAGGTTGAAGGAGAGATCTTGCAAGGTGTTCTTACCAGATTGTTTTCGAGAGACATATTGTTGAAAT  
GCAAAATCTACTCCGACGTTCTTTGGGACAAGAAGTACTCAGCTTGAAACCAATGCGGTTTTATTGAAGATTTATTG  
GCAGAGCACCTGTTGGTGATGAAGATAATTAA

## 4256.2 (SEQ. ID. NO. 359)

ATGAAAAACAAGCCTATGTCAATTATTGCTCTCACCTCCTCTCTATTTGTCTTTTTTTCTCCACAGCTTGCTGGAAATACTT  
GATTTTGACTGGTCTATCTTTTTGCACGATGTCGAAAAACAGAAAAATTTGTCTTTTTATTGTTGGTTTTACAGCATGTCCATG  
ACCTGTCTCTTAGCCCTGTTTTGGCGAGGGATCGAAGAGCTTTCTCTAAGAAAAATGCAGGCTAATCTCAAGCGTTTATTAGC  
AGGGCAAGAAGTGGTTACAGGTTGCAGATCCAGATTGGATGCCAGTTTCAAGTCCTTATCAGGTAACTTAACTTTTGACA  
GAGGCTCTTCAAAAAGCTGAAAAATCAGAGCCTTGCTCAGGAAGAGGAAATCATCGAGAAGGAACGGAAGCGAATTGCTCGG  
GATTTGCACGATACAGTCAGTCAGGAGTTGTTTGGCGCCACATGATTTTATCGGGTATCAGTCAGCAGGCTTTGAAATTGGA  
TAGAGAAAAGATGCAGACCCAGTTGCAGAGTGTACAGCTATTTAGAAACAGCCAGAAAGGATTTGCGGGTTTTGCTCTTG  
CATTTGCGACCAAGTTGAACTGGAGCAGAAGAGCTTGATAGAAGGGATTCAAATCTTTTAAAGAGCTTGAGGACAAGAGTG  
ATCTTAGGGTTAGTCTCAAGCAGAATATGACGAAATGGCTAAGAAAAATCGAGGAGCATATCTTCCGTATCCTGCAAGAGTT  
GATTAGCAATACCCTCCGCCATGCCAGGATCTTGCTAGATGTTCTACCTCTATCAGACAGATGTTGAATTGCAATGGAAGG  
TGGTGGACAATGGGATTGGTTCCAGTTAGGGAGCTTAGACGACTTGAGTTATGGACTGCGAAATATCAAGGAGCGGGTTGA  
AGATATGGCTGGAACAGTTCAACTCTTGACAGTCCCAAGCAAGGGCTGGCGGTTGATATCCGTATTCCTCTGTTAGATAAG  
GAATGA

## 4263.1 (SEQ. ID. NO. 360)

ATGATGTTTCCATTATTTCTCAAGGATTTGTCTGGGCTATTCTAGGTCTGGGAATCTTTATGACATTTAGGATTTTAACTTT  
CCAGATATGACGACAGGAAGGTTCCCTCTTGGGGGAGCTGTTGCTGTCACTTTGATAACCAAGGCGTGAACCCATTTT  
AGCGACACTTGTGCTGTAGGAGCAGGTTGTTTGGCTGGAATGGCAGCAGGCCCTCTTTATACAAAAGGGAAGATCCCAACC  
TTGCTCTCAGGGATTTTGGTGATGACTTCTGTCACTCAATCATGCTCTTGATTATGGGACGTGCCAATTTAGGCCTGCTTGA



ACCAAGCAAATTCAGGATGTTTTGCCCTTTTGATTCCGGATTTGAATCAACTCTTGACAGGTCTCATCTTTGTGAGTATTGTTATT  
GCTCTCATGCTCTTTTTCTTGGACACTAAACTCGGACAAGCCTATATTGCTACAGGGGATAATCCTGATATGGCTAGAAGTTT  
CGGGATTCACTACTGGACGCATGGAGCTCATGGGCTTGGTCTTATCAAATGGTGTGATTGCCCTTGACAGGTGCCCTCATTGCTC  
AGCAAGAAGGTTATGCCGATGTGTCTCGAGGGATCGGGGTTATCGTTGTGGGGCTTGCAAGTTTGATTATTGGAGAAGTTAT  
TTTCAAGAGTTTGAGCTTTGGCAGAGCGTTTGGTTACTATCGTTGTAGGTTCTATCGCTTATCAATTTTTAGTGTGGGCAGTTAT  
CGCATTGGCTTTAATACAAAGTTACCTTCGTTTATACAGTGCCTTGATTTTAGCAGTCTGCCTCATGATTCCAACATTTAAGCA  
AACAATCTTGAAAGGAGCCAAGTTAAGCAAATGA

## 4346.1 (SEQ. ID. NO. 361)

ATGAAAAAATGAAAGTTTGGTCTACTGTACTTGCAACGGGAGTTGCTCTTACTACACTTGCTGCTTGCTCTGGAGGTTCAAA  
TTCTACGACTGCTTCTTCATCTGAAGAAAAAGCTGATAAAAGTCAAGAATTAGTTATCTATTTCGAAGTCAAGTCTCAAAATGGTC  
GTGGTGATTGGTTAACTGCTAAAGCAAAAGAAGCTGGTTTAAATATAAAAAATGGTTGATATCGCTGGCGCTCAATTAGCAGA  
CCGTGTTATTGCTGAGAAGAAATATGCAGTTGCAGATATGGTATTGGAAATGGTGCTGTTGATTCAAATAAAATAGAGATC  
AAAAATTACTAGTACAGTACAAGCCTAAATGGTTAGATAAAATTGATCAATCTTATCAGATAAAGATAATTATTATAATCCT  
GTGATTGTTCACCATTAAGTTTAAATTTGGGGCGCCTGATGTAAAAGAAATGCCTAAAGATTGGACTGAATTAGGTAGTAAGT  
ATAAAGGTAAATATTCAATTTCTGGTCTTCAAGGAGGTACAGGACGGGCAATTCTAGCAAGTATCTAGTTCGATACCTTGAT  
GATAAAGGTGAATTAGGTGTTTCCGAAAAAGGTTGGGAAGTAGCAAAAAGAAATATTTGAAAAATGCATACACTCTTCAAAAG  
GGAGAAAGTTCAATTGTTAAGATGTTAGACAAAAGAAGATCCAATACAATATGGAATGATGTGGGGTCTGGTGCATTAGTTG  
GACAAAAAGAACAAAATGTTGTTTCAAAGTTATGACTCCTGAGATTGGTGTACCATTTGTAAGTGAACAACTATGGTTTAA  
AGCACTAGTAAAAAACAAGCGTTAGCTAAAGAAATTTATTGATTGGTTGGTCAATCAGAAATTCAGTGAATATAGTAAGA  
ACTTTGGATCTATTCTGCAATAAAGATGCCCTCAAAGATCTACCTGAAGATACGAAGAAATTTGTTGATCAAGTGAACCC  
ACAAAATATTGACTGGGAAGCTGTTGGAAAGCATTGGATGAATGGGTAGAAAAAGCTGAATTAGAATACGTACAATAA

## 4346.2 (SEQ. ID. NO. 362)

ATGATTAATTTGATAATATTCAAATTAATATGGTGATTTTGTGCAATTGATAATCTGAATTTAGATATACATGAAGGGGA  
ATTTTTTACATTTCTTGGGCTTCAGGATGTGGTAAATCAACTACTTTGAGAGCATTGGTAGGTTTTCTAGATCCATCATCAGG  
AAGTATTGAAGTTAATGGAACAGATGTCACCTCATTTGGAACCTGAAAAGCGTGGAATTGGTATTGTAATTTCAATCTTAGCGC  
TATTTCCAACATGACTGTTTTTGTATAATATTGCATTTGGTTTAAAAGTTAAGAAGGTAGCTCCAGATGTTATTAAGCTAAA  
GTATCAGCAGTGGCAGCAAAAATTAAGATCTCTGATCAACAGTTACAGCGTAATGTATCAGAATTATCTGGGGGTCAACAAC  
AAAGGGTAGCATTGGCTCGTGTCTGTTCTTGAACCTAAAATTCCTTGTCTAGATGAACCATTTGTCAAACCTTGACGCAAAA  
TTACGTGTAGATTGAGAAAAGAGTTGAAAAGACTTCAAAAAGAGTTAGGTATTACTACTTTATATGTTACTCATGATCAAG  
AGGAAGCCTTGACTTTATCTGATAGAATTGCAGTCTTAAACAATGGATACATCGAACAGGTGGTACACCAGTAGAGATTTA  
TCATAATTCTCAAACCTGAATTTGTATGTGATTTTATTGGAGATATTAATGTTTGGACCGATGAAACAGTCCACGAAGTATTAT  
TGAAAAATACAAAGCGTTTTCTTAGAGGATAAAAAAGGATACATTCGATTAGAGAAAGTTCGATTCAATCGTGAACCTGAACA  
AGATTTTATTCTAAAAGGGACAATTATTGATGTTGAGTTTTCTGGAGTTACAATTCACTATACAATAAAAGTTTCTGAAAGTC  
AGATTCTTAATGTAACAAGTATTGATAGTCAGGCTGCTATTAGATCTGTGCGAGAAAGTGTGGAATTATTTATCACACCATCA  
GACGTTCTGCAATTTTAA

## 4346.3 (SEQ. ID. NO. 363)

ATGCGTCATAAATTAATTTAAAAAGATTGGCTTATTCGTTTAGGGTTAATCTGGTCTTAGTAACATTTATTATTTATCCAAAC  
TTTGATCTAGTAGTGAATGTATTTGTAAGAGGAGGAGAAATTTCCCTTGATGCTGTACATCGTGTCTAAAAATCTCAGAGGGC  
ACTTCAGAGTATTATGAACAGTTTTAAGTTAGCATTTTCACTCATTATTACAGTTAATGTCGTAGGTATTCTTTGTGTTCTATT  
TACAGAGTACTTTGATATTAAAGGTGCTAAAATTTTAAAAATTAGGTTATATGACCTCTTAAATTTATGGAGGAGTGGTTTTAG  
CGACTGGATATAAATTTGCTATGGTCCTTATGGATTGATTACAAAAATTTTACAAAATGTTATCCCTTCTTTAGACCCTAACT  
GGTTTATTGGGTATGGTGCAGTCTTATTCATTATGACATTTTCAGGAACCTGCTAATCATACTGTTTAAAAACAAATACAATTC  
GAAGCGTTGACTATCACACTATTGAGGCTGCTCGAAATATGGGAGCAAAACCATTTACTGTTTCCGAAAAGTAGTGTACC  
AACCTTAATTTCAACTCTATTTGCACTTACTATTAATGGTTTTCTTAGTGGTTTATCTGCAGTAGCAGCACCCATGATTGTTGG  
TGTTAAAGAAATTTCAAACATAAATCCAATGATTATTACATTTGCAGGGATGGGGAATTCTCGTGATTAGCTGCCCTACTTG  
CAATTTATTTAGGTATTGCAACTACAATTTTGCTTACTATCATGAATAAGATAGAAAAAGGTGGAAATTATATTTCTATCTCT  
AAGACTAAAGCGCCTCTTAAAAAACAAAAATTCGCTCTAAGCCTTGGAATATCATTTGCTCACATTGTAGCATATGGATTGT  
TCACAGTTTTTATGCTTCCACTAATTTTTATAGTATTATACTCATTACAGATCCAGTTGCAATTCAAACAGGTAACTTAACAT  
TATCAAACCTTACTTTAGAAAAATTATCGCTTATTCTTTAGTAATAGTGCAGGATTTCTCTCCATTCTTGGTCAAGCTTTATTATTTC  
TATTATTGCTGCGACAACAGCAACAATTTCTCGCAGTTGTATTGCTCGTGTGTCAGAAAACATAAATCTCGTTTGTATTCTT  
ATTTGAATATGGTGCTTACTTCTTGGTTACTACCAAGTACACTTTTAGCAGTAAGTTTATTATTTACTTTTAAATCAGCCACA  
ATTTCTTGTCTTGAATCAGATTTTGGTAGGTAGTTTGGTAATTTACTATTGTCATATAGTTGTAATAAAATCCCATTTTCTTAT  
AGAATGGTACGTGCTATTTTATTTAGTGTGATGATGAGATGGAAGATGCAGCAAGAAGTATGGGTGCTTACCCTTTTATAC  
TATGATGAAGGTTATCATTCCATTTATTTACCGGTTGTTCTCTCTGTTATTGCTTTAAACTTTAACTCTTTTAACTGACTTC  
GACTTATCTGTATTTCTTTACCATCCCCTAGCTCAACCATTAGGTATTACGATTGATCTGCAGGTGATGAAACAGCAACATC  
TAATGCACAAGCTCTGGTATTTGTTTATACAATTTGTTCTGATGATTATTTCTGGAACGGTATTATACTTCACACAAGACCGG  
GGCGTAAAGTAAGGAAATAA



Table 2

(SEQ. ID. NO. 1)

MEELVTLDCFLIDRTKIEANANKYSFVWKKTTTEKFSAKLQEQIQVYFQEEITPLLIKYAMFDKKQKRGYKESAKNLANWHYNDKE  
DSYTHPDGWYRYRHHHTKYQKTQDFQEQEIKVYYADEPESAPQKGLYMNERYQNLKAKECQALLSPQGRQIFAQRKIDVEPVFGQI  
KASLGKRCNLRGKRQVRIDMGLVLMANNLLKYSKMKZ

(SEQ. ID. NO. 2)

MGKGHWNRKRVSIRKFAVGACSVMIGTCAVLLGGNIAGESVUYADETLTHTAEKPKEEKMIVEEKADKALETKNIVERTEQSE  
PSSTEAIASEKKEDEAVTPKEEKVSAKPEEKAPRIESQASNQEKPLKEDAKAVTNEEVNQMIEDRKVDNFQNWYFKLNANSKEAIK  
PDADVSTWKKLDLPYDWSIFNDFDHESPAQNEGGQLNGGEAWYRKTFKLDEKDLKKNVRLTFDGVYMDSDQVYVNGQLVGHYP  
NGYNQFSYDITKYLQKDGRENVIHVAHVNKQPSSRWYSGSGIYRDVTLQVTDKVHVEKNGTTILTPKLEEQQHKGKVVETHVTSKIV  
NTDDKDHELVAEYQIVERGGHVAVTGLVRTASRTLKAHESTSLDAILEVERPKLWTVLNDKPALYELITRVYRDGQLVDAKKDLFG  
YRYHHWTPNEGFSLNGERIKFHGVSLLHHDHGAALGAEENYKAEYRRLKQMKEMGVNSIRTTNHPASEQTLQIAAELGLLVQEEAF  
DTWYGGKKPYDYGRFFEKDATHEARKGEKWSDFDLRTMVERGKNNPAIFMWSIGNEIGEANGDAHSLATVKRLVKVIKDVVK  
TRYVTMGADKFRFGNSGGHEKIADELDAVGFNYSEDNYKALRAKHPKWLIYGETSSATRTRGSYIRPERELKHSNGPERNYE  
QSDYGNDRVGWGTATASWTFDRDNAGYAGQFIWTGTDYIGEPTPWHNQNQTPVKSSYFGIVDTAGIPKHDFLYQSQWVSVK  
KKPMVHLLPHWNWENKELASKVADSEGKIPVRAYSNASSVELFLNGKSLGLKTFNKKQTSQDGRTYQEGANANELYLEWKVAYQ  
PGTLEAIARDESKEIARDKITTAGKPAAVRLIKEDHAIADGKDLTYIYIEIVDSQGNVPTANNLVRFLHGGQQLVGVNDGEQ  
ASRERYKAQADGSGWIRKAFNGKGVAVIKSTEQAGKFTLTAHSDLLKSNQVTVFTGKKEGQEKTVLGTVEVPKVQTHIEAPEMPTT  
VPFVYSYDGSRAERPVTWSSVDVSKPGIVTVKGMADGREVEARVEVIALKSELPPVKRIAPNTDLNSVDKSVSYVLIDGSVEEYEV  
KWEIAEEDKAKLAIPGSRIGATGYLEGQPIHATLVVEEGNPAAPAVPTVTVGGEAVTGLTSQKPMQYRTLAYGAKLPEVTASAKN  
AAVTVLQASAAANGMRASFIQPKDGGPLQTYAIQFLEEAPKIAHLSLQVEKADSLKEDQTVKLSVRAHYQDGTQAVLPADKVTFS  
TSGEVEAIRKMGLELHKGPAVTLNAEYEGAKDQVELTIQANTEKKIAQSIRPNNVVDLHQPSPATVTVYEDKGFPPKTHKVT  
WQAIPKEKLDYSYQTFEVLGKVEGIDLEAKVSVSEGVSVVEVSVTTPIAEAPQLPESVRTYDSNGHVSSAKVAWDAIRPEQYAKE  
GVFTVNGRLEGTQLTKLHVRVSAQTEQGANISDQWTGSELPLAFASDSNPSDPVSNVNDKLSYNNQPANRWTNWNRNTPNEASV  
GVLFGDSGILSKRSVDNLSVGFHEDHGVGPKSYVIEYYVGKTVPTAPKNPSFVGNEDHVFNDNSANWKPVTNLKAPAQLKAGEM  
NHFSFDKVETAVRIRMVKADNKRGTSTIEVQIFAKQVAAAKQGQTRIQVDGKDLANFNPDLDYDYESVDGKVPVAVTASVSN  
GLATVVPVSVREGEPRVIAKAENGIDILGEYRLHFTKDKSLSHKPVAAVKQARLLQVGGQALELPTKVPVYFTGKDGKETKDLTVE  
WEEVPAENLTAKAGQFTVRGRVLGSNLVAEITVRVTDKLGELSDNPNYDENSNAFASATNDIDKNSHDRVYDLYNDGDHSENRR  
WTNWSPTSSNPESVAGVIFRENGKIVERTVTVQKGVFFADSGTDAPSRLVLERVVGPEFEVPTYYSNYQAYDADHPFNPNPENWE  
AVPYRADKDIAAGDEINVTFAIKAKAMRWMERKADKSGVAMIEMTFLAPSELPESTQSKILVDGKELADFAENRQDYQITYK  
GQRPKVSVEENNQVASTVVDSDGSDSPVLVRLVSESGKQVKEYRIHLTKEKPVSEKTVAAVQEDLPKIEFVEKDLAYKTVEKKDS  
TLYLGETRVEQEGKVGERIFTAINPDGSKKEKLREVVPTDRIVLVGTPVAQEAKKPVSEKADTKPIDSSSEASQTNKAQLPST  
GSAASQAAVAAGLTLGLSAGLVVTKGKKEDZ

(SEQ. ID. NO. 3)

MKIMKKKYWTLLAILFFCLFNNSVTAQEIPKNLDGNITHTQTSSESSESDEKQVDYSNKNQEEVDQNKFRIDKTELFTVTDKHLK  
NCKLELEPQINNDIVNSESNNLLGEDNLDNKIKENVSHLDNRGGNIEHDKDNLESSIVRKYEWIDDKVTGGGESYKLYSKSNSKV  
SIALIDSGVDLQNTGLLKNLSNHSKNYPNKGYLKGEEGEGEHSIDQDRLGHTAVVAQIVGDDNINGVNPVHNINVRIFGKSSA  
SPDWIKAFIDVADDDGNDIINLTGQYLMIDGEYEDGTNDFETLKYKAIDYANQKGVIIAALGNDLSNDSKNEARKVIAQKRRK  
VRKPLGVVDVPSYFSSTISVGGIDRLGNLSDFSNKGSDAIYAPAGSTLSLSELGLNNFINAEKYKEDWIFSATLGGYTYLYGNSFA  
APKVSQAIAIMIIDKYKLDQPYNYMFVKKFWKKHYQZ

(SEQ. ID. NO. 4)

MKKTWKVFLTLVTALVAVVLVACGQGTASKDNKEAELKKVDFILDWTPNTNHTGLYVAKEKGYFKEAGVDVDLKLPPPESSSD  
LVINGKAPFAVYQDYMAKKLEKGAGITAVAAIVEHNSTSGIIRKSDNVSSPKDLVGKKYGTWNPTELAMKLTLVESQGGDFEK  
VEKPNNDNSITPIANGVFDTAWIYYGWDGILAKSQGVANFMYLKDYYKEFDYSPVIANNDYLDKNKEARKVIAQKRRK  
QYAMEHPPEEAADILIKNAPELKEKRDVFIESQKYSKEYASDKEKWGQFDAARWNAFYKWDKENGILKEDLTDKGFTNEFVKZ

(SEQ. ID. NO. 5)

MKRTWRNSFVTNLNTPFMIGNIEIPNRTVLAPMAGVTNSAFRTIAKELGAGLVVMMEMVSDKGIQYNNKTLHMLHIDEENPVSI  
QLFGSDEDSLARAEEFIQENTKTDIVDINMGCPVNKIVKNEAGAMWLKDPDKIYSINKVQSVLDIPLTVKMRGTWADPSLAVENA  
LAAEAAGVSALAMHGRTRQMYTGHADLETLYKVAQALTKIPFIANGDIRTVQEAQRIEEVGADAVMIGRAAMGNPYLFNQIN  
HYFETGEILPDLTFEDKMKIAYEHLKRLINLKGENVAVREFRGLAPHYLRGTSGAAKLRGAISQASTLAEIETLLQLEKAZ

(SEQ. ID. NO. 6)

MIKNPKLLTSKFLRSFAILGGVGLVIHIAIYLTFFPYYIQLEGEKFNESARVFTEYLKTKTSDEIPSLQSYSKSLTISAHLKRDIVDKR  
LPLVHDLIDKDGKLSNYIVMLDMSVSTADGKQVTVQFVHGVDVYKEAKNILLLYLPYTLVTFIAFSFVFSYFYTKRLLNPLFYISEV  
TSKMQLDDNIRFDESARKDEVGEVGKQINGMYEHLKVIYELESRNEQIVKLQNKQVSFVRGASHELKTPLASRLRIENMQHNIG  
DYKDHPKYIAKSINKIDQMSHLLLEEVLESSKFQEWTECRETTVPVLVDILSRYQELAHSIGVTIENQLTDATRVMMSLRALDKVLT  
NLISNAIKYSKDNKGRVISEQDGYLSIKNTCAPLSDQELEHLDFIFYHSQIVTDKDESSGLGLYIVNNILESYQMDYSFLPYEHGMEFK  
ISLZ

(SEQ. ID. NO. 7)

MYLGDLMEKAECGQFSILSFLQESQTTVKAVMEETGFSKATLTKYVTLNDKALDSGLELAHSEDENLRLSIGAATKGRDIRSLF  
LESVYKYQILVYLLYHQFLAHQLAQELVISEATLGRHLAGLNQILSEFDLSIQNGRWGPEHQIHYFYFCLFRKVWSSQEWEGHM  
QKPERKQEIANLEEICGASLSAGQKLDLVLAHISQQLRLVNACQFQVIEEKMRGYFDNIFYLRLLRKVPSSFAGQHIPLGVEDGE  
MMIFFSFLLSHRILPHTMEYILGFGGQLADLLTQLIQEMKEELLGDTYEDHVTYELSQLCAQVYLYKGYILQDRYKYQLENRHP  
YLLMEHDFKETAEELFIHALPAFQQTDLDDKILWEWLQLEIYMAENGQQHMRIGDLTSGFLVFSRMAAILKRYLEYNRFITIEAY  
DPSRHYDLLVTNNPIHKKEQTPVYYLKNLDLMDLVAIRQLLFTZ

(SEQ. ID. NO. 8)

MEFSKKTRELSIKKMQERTLDLLIIGGGITGAGVALQAAASGLETLGIEMQDFAEGTSSRSTKLHGGGLRYLKQFDEVVSDTVSE  
RAVVQIAPHIPKSDPMLLPVYDEDGATFSLFRLKVAMDLYDLLAGVSNTPAANKVLSKDQVLERQPNLKKEGLVGGGVYLDLFR

NNDARLVNIENIKRANQD GALIANHVKAEGFLFDES GKITGVVARDLLTDQVFEIKARLVINTTGPWSDKVRNLSNKGTFQFSQMRPT  
KGVHLVVDSSKIKVSPQVYFD TGLGDGRMVFLPRENKTYFGTTDDTYTGDLEHPKVTQEDVDYLLGIVNNRFPESNITIDDISS  
WAGRLPLIAGNSASDYNGGNGTISDESFDNLIA TVESYLSKEKTREDVESAVSKLESSTSEKHLDPASVSRGSSLD RDDNGLLTLA  
GGKITDYRKMAEGAMERVV DILKAEFDRSFKLINSKTYPVSGGELNPANVDSEIEAFAQLGVSRGLDSKEAHYLANLYGSNAPKV  
FALAHSLEQAPGLSLADTSLHYAMRNELTSPVDLLRRTNHMLFMRDSLDSIVEPILDEMGRFYDWTEEEKATYRADVEAALA  
NNDLAE LKNZ

(SEQ. ID. NO. 9)

MMNELFGEFLGTLILILLGNVGVAGVVL PKTKSNSSGWIVITMGWGIAVAVAVFVSGKLPAYLNP AVTIGV ALKGGLPWASVLP  
YILAQFAGAMLGQILVWLQFKPHYEAENAGNILATFSTGPAIKD TVSNLISEILGT FVLVLTIFALGLYDFQAGIGTFAVGTLIVGIG  
LSLGGTTGYALNPARDLGPIMHSILPIPKNKG DGDWSYAWIPVVG PVIGAALAVLVFSLFZ

(SEQ. ID. NO. 10)

MTKKKIERISVIHREKILWLKWFYMRDKEQPKYSVLERKMFDAAKNQDMLAYQKYATIKQITDIRVQTSEADILEAVKEVYVYNH  
MNVIGACQRLFISQSPAYDKLNKWFNIYSDLYFSVPLPKMGVYHEMVGIZ

(SEQ. ID. NO. 11)

MKNSNEAEMKLLYTDIRTSLEILTREAELVAAGKRVFYIAPNSLSFEKERA VLEYLSQQA SFSITVTRFAQMARYLV LNDLPAKT  
TLDDIGLGLAFYKCLAELDPKDLRVYGAIKQDPQLIQQLIELYHEMTKSQMSFLDLENLTDEDK RADLLLIFEKVTA YLNNQGLAQ  
ESQLSHLIEAIENDKVSSDFNQIALVIDGTRFS AEEERVVDDLHGKGVEIVIGAYASKKAYTSPFSEGNLYQASVKFLHHLASKYQT  
PAQDCSQTHEKMDSFDKASRLLESSYDFSELALDVDEKDRENQIWSCLTQKEELVARSIRQKLHENS DLSYKHFRILLGDVAS  
YQLSLKTIFDQYQIPFYLRSEAMAHPLTQFVESILALKRYRFRQEDLINLLRTDLYTDLSQSDIDAFEQYIRYLGINGLPAFQQTFT  
KSHHGKFNLERLNLRLRILAPLETLFASRKQKA ELLQKWSVFLKEGAVTKQLQDLTTTLEAVEQERQAEVWKAFCHVLEQFAT  
VFAGSQVSLEDFLALLHSGMSLSQYRTIPATVDTVLVQSYDLIAPLTADFVY AIGLTQDNLPKISQNTSLTDEERQNLNPAETEEGV  
QLLIASSENKKNRYTMLS LVSARKQLFLSAPSLFNESESKE S AYLQELIHFGFRREKRMNHKGLSKEDMGSYHSLSSLVAYH  
QQGEMSDTEQDLTFVKVLSRVIGKLDQQGLENP IPTSPSKTLAKDTLQALYPAKQEFYLS TSGLTEFYRNEYSYFLRYVLGLQ  
EELRLHGNPDARLHNRFLHRIFERALQLPNEDSFDQRLEQA IQETSQEREFEAITYQESLEAQFTKEVLLDVARTTG HILRHNP AETIKEE  
ANFGGKDQAFIQLDNGRSV FVRGKVDRIDRLKANGAIGVVDYKSSLTQFQFPFFHFNGLNSQLPTYLAALKREGEQNFFGAMYLEM  
AEPVQSLMAVKSLAGAVVEASKSMKYQGLFLEKESYLGFEFYNKNKANQLTDEEFQLLLDYNA YLYKKA AEKILAGRFAINPYT  
ENGRSIAPYVQHQAITGFEANYHLGQARFLEKLDLADGKRLVGEK LKQAWLEKIREELNRZ

(SEQ. ID. NO. 12)

MKLIPFLSEEEIQLQEA EANSSKEQKKTAEQIEAITYSAQNILVSASAGSGKTFVMAERILDQLARGVEISQLFISTFTVKAATELKE  
RLEKKISKKIQETDDVDLQHLGRQLADLPNAAIGTMSDFTQKFLGKHGYLLDIAPNFRILQNQSEQLILENEVFHEVFEAHYQ GK  
QKETFSHLLKNFAGRGKDERGLRQQVYKIYDFLQSTSNPQKWLSESLKGF EKADFTSEKEKLTEQIKQALWDLESFFRYHLDND  
AKEFAKAA YLENVQLILDEIGSLNQESDSQAYQAVLARVVAISKEKNGRALTNASRKADLKPLADAYNEERKTQFAKLQGLSDQI  
AILDYQERYHGD TWKLAKTFQSFMSDFVEAYRQRKRQENAFEFADISHYTIEILENFPQVRESYQERFHEVMVDEYQDTNHIQER  
MLELLSNGHNRFMVGD IKQSIYRFRQADPQIFNEKFQRYAQNPQEGRLIILKENFRSSSEVLSATNDVFERLMDQEVGEINYDNKH  
QLVFANTKLT PNPNDKAAFLYDKDDTGE EESQTETKLTGEMRLVIKEILKLHQEKGVAFKEIALLTSSSRNDQILLALSEYGIP  
VKT DGEQNNYLSQSLVQVMLDTRLVIHNP LQDYALVALMKSPMFGFDEDELARLSLQKAEDKVHENLYEKLVAQKMASSQKG  
LIHTLAELKQFMDILASWRLYAKTHSLYDLIWKIYNDRFYDYDVGALPNGPARGANLYALALRADQFEKSNFKGLSRFIRMID  
QVLEAQHDLASVAVAPPKDAVELMTIHKSKGLEFPYVFILNMDQDFNKQDSMSEVILSRQNGLG VKYIAKMETGAVEDHYPKTIK  
LSIPS LTYRQNEELQLASYSEQMRLLYVAMTRA EKKLYLVGKGSREKLESKEYPAAKNGKLSNTRLQARNFQDWLWAIKSVFT  
KDKLNFYRFIGEDQLTREAI GELETKSPLODSSQADNRQSDTIKEALEMLKEVEVYNTLHRAAIELPSVQTPSQIKKFYEPVMDME  
GVEIAGQGGSVGGKISFDLPDFSTKEKVTGAEIGSATHELMQRIDLSQQLTLASLTETLQVQVTSQAVRDKINLDKILAFFD TVLGQ  
EILANTDHL YREQPFSMLKRDQKSQEDFVVRGILDGYLLYENKIVLFDYKTD RYDEPSQLVD RYRGQALALYEEALS RAYS IENIEK  
YLILLGKDEVQVVKVZ

(SEQ. ID. NO. 13)

MELARHAESLGVDIAITIPPIYFRLPEYSVAKYWN DISSAAPNTDYVIYNIPQLAGVALT PSLYTEMLKNPRVIGVKNSSMPVQDIQ  
TFVSLGGEDHIVFNGPDEQFLGGRLMGARAGIGGTYGAMPELFLKLNQLIADK DLETARELQYAINAIIGKLTSAHG NMYGVIKEV  
LKINEGLNIGSVRSPLTPVTEEDRPVVEAAAALIRETKERFLZ

(SEQ. ID. NO. 14)

MYKTKCLREKLVFLKIFFPILYQFANYSASFVDTAMTGQYNTMDLAGVSMATSIWNPFFTFLTGTIVSALVPIIHHHLGRGKKEEV  
ASDFYQFIYALGLSVLLGMVFLAPILNHIGLEAAVA AVRYLWFLSIGIPLLLFSVIRSLLDSLGLTKLSMYLMLLLPLNSG  
FNYLLIYGAFGVPELGGAGAGLGTSLAYWVLLGISVLVLFKQEK LKALHLEKRIPLNMDKIKEGVRLGLPIGGTVFAEVAIFS VVGL  
IMAKFSP LIIASHQSAMNFSSLMYAFPMSSISAMAI VVSVEVGAKRFD DAKTYIGLGRWTALIFA AFTLTFLYIFRGNVASLYGN DP  
KFIDLTVRFLTYSLFFQLADTFAAPLQGILRGYKDTVIPFYLGLLGYWGVAIPVYAI Z

(SEQ. ID. NO. 15)

MSTLAKIEALLFVAGEDGIRVRQLAELLSLPPTGIQQSLGKLAQKYEKDPDSSLAIETSGAYRLVTKPQFAEILKEYSKAPINQSLS  
RAALETLSIIAYKQPI TRIEIDAIRGVNSSGALAKLQAFDLIKEDGKKEVLGRPNLYVT TDYFLDYMGINHLEELPVIDELEIQAQESQ  
LFGERIEEDENQZ

(SEQ. ID. NO. 16)

MDTMISRFRHLFEAL KSLKRNGWMTVA AVSSVMITLTLVAIFASVIFNTAKLATDIENNVRVVVYIRKDVEDNSQTIEKEGQTVT  
NNDYHKVYDSLKNMSTVKS VTFSSKEEQYEKLTEIMGDNWKIFEGDANPLYDAYIVEANTPN DVKTIAEDAKKIEGVSEVQDGG  
ANTERLFLKASFIRVWGLGIAALLIFIAVFLISNTRITISRSREIQIMRLVGAKNSYIRGPFLLEGAFIGLLGAIAPSVLVFIVYQIVYQS  
VNKSLVGQNLMSIPDLFSPLMIALLFVIGVFIGSLGSGISMRRFLKIZ

(SEQ. ID. NO. 17)

MKKVRFIFLALLFFLASPEGAMASDGTWQKGQYLKEDGSQAANEWVFDTHYQSWFYIKADANYAENEWLKQGDDYFYLSKSGG  
YMAKSEWVEDKGAFYLDQDQGMKRNAWVGTSYVVGATGAKVIEDWVYDSQYDAWIFYIKADGQHAKEWLOQIKGKDYFYKS  
GGYLLTSQWINQAYVNASGAKVQQGWLFQKQYQSWFYIKENGNYADKEWIFENGHYFYLSKSGGYMAANEWIWDKESWIFYLK  
FDGKMAEKWVYDSSHQA WYFYKSGGYMTANEWIWDKESWIFYLKSDGKIAEKWVYDSSHQA WYFYKSGGYMTANEWIWD  
KESWIFYLKSDGKIAEKWVYDSSHQA WYFYKSGGYMAKNETVDGYQLGSDGKWLGGKTTNENAA YQVVPVTANVYDSDGE  
KLSYISQGSVWLDDKDRKSDDKRLAITISGLSGYMKTEDLQALDASKDFIPYYESDGHRYHYVAQNASIPVASHLSDMEVGKKY  
YSADGLHFDGFKLENPFLFKDLTEATNYSAEELDKVFSLLNNNSLLENKGATFKEAEHYHINALYLLAHSALSNWGRSKIAKD  
KNNFFGITAYDTTPYLSAKTFDDVDKGILGATKWIKENYIDRGRITFLGNKASGMNVEYASDPYWGEEKIASVMMKINEKLGGKZ

(SEQ. ID. NO. 18)

MKKVLQKYWAWAFVVIPLLLQAIFFYVPMFQGAFFYSFTNWTGLTYNYKFVGLNNFKLLFMDPKFMNAIGFTAIIAIA MVVGEIAL  
GIFIARVLNSKIKGQTFRAWFFFPVLSGLTVALIFKQVFNYGLPAIGNALHIEFFQTSLLGTKWGAIFA AVFVLLWQGVAMPIIIFL  
AGLQSIPTTEAARIDGATSKQVFWNIELPYLLPSVSMVFILALKGGLTAFDQVFAMTGGGPNNATTSLGLLVNYAFKNNQFGY  
ANAIAVILFLLIVISIIQLRVSKKFEIZ

(SEQ. ID. NO. 19)

MMKQDERKALIGKYILLILGSVLILVPLLATLFSFPTKDIVDNFFGFPTNFTWDNFSRLLADGIGGYWNSVVTIVLSLLAVMIFIP  
MAAYSIARNMSKRKAFTIMYTLLILGIFVPFQVIMIPITVMMSKLGLANTFGLILLYLTAYIPQTLFLYVGYIKISIPESLDEAAEIDGA  
NQFTTYFRIIFPMMKPMHATTMIINALWFWNDFMLPLLVLNRDSKMWTLPFLQYNYAGQYFNDYGPSFASYVVGIIISITIVYLFQ  
RHIISGMSNGAVKZ

(SEQ. ID. NO. 20)

MKSILQKMGHEPMLLLFLSYSTVISILAQNWMLVASVGMFLFTIFFLHYQSILSHKFFRLILQFVLFGSVLSAASFASLEHFQIVKKF  
NYAFLSPNMQVWHQNRAEVTFNPNYYGIICFCMIAFYLFITTKLNLWKVFCVIAGFVNLFGNFTQNRATAPAIAGAIYLFITTI  
KNWKAFWLSIGVFAIGLSFLFSSDLGVRMGTLDSSMEERISIW DAGMALFKQNPFWGEGPLTYMNSYPRIHAPYHEHAHSYIDITI  
LSYGIVGTILLVSSVAPVRLMMDMSQESGKRPIIGLYLSFLTVAHVHGFIDLALFWIQSGFIFLLVMCSIPLEHRLVSDMTDZ

(SEQ. ID. NO. 21)

MSKMDVQKIIAPMMKFVNMRGIIALDKGMLAILPLTVVGSFLIMGQLPFEGNLKSIASVFGANWTEPFMQVYSGTFAIMGLISCF  
IAYS AKNSGVEALPAGVLSVSAFFILLRSSYIPKQGEAIGDAISKVWFGGQGIAGIIIGLVVGSYITFFIKRKIVIKMPEQVPQAIKQ  
FEAMIPAFVIFLSSMIVYILAKSLTNGGTFIEMIYSAIQVPLQGLTGSLYGAIGIAFFISFLWWFGVHGQSVVNGVVTALLSNLDAN  
KAMLASANLSLENGAHIVTQQFLDSFLILSGSGITFGLVVAAMLFAAKSKQYQALGKVAAPFAIFNVNEPVVFGFPIVMNPVMFVPFI  
LVPVLA AVIVYGA IATGFMQPFSGVTLPWSTPAILSGFLVGGWQGVITQLVILAMSTLVYFPFFKVQDRLAYQNEIKQSZ

(SEQ. ID. NO. 22)

MKKKDLVDQLVSEIETGKVRTLGIYGHGASGKSTFAQELYQALDSTTVNLETPYITSGRHLVVPKDAPNQKVTASLPVAHELES  
LQRDILACRRVWMSZ

(SEQ. ID. NO. 23)

MKKRYLVLTALLALSAAQSQEKTKNEDGETKTEQTAKADGTVGSKSQGAQAQKAEVNVNKGDYYSIQGKYDEIIVANKHYPLSK  
DYNPGENPTAKAELVKLIKAMQEAGFPISDHYSGFRSYETQTKLYQDYVNQDGKAAADRY SARPGYSEHQTGLAFDVIGTDGDL  
VTEEKAAQWLLDHAADYGFVVRYLKGKEKETGYMAEEWHLRYVGKEAKEIAASGLSLEEYGFEGGDYVDZ

(SEQ. ID. NO. 24)

MREPDLNHLKKGKGYFKKHAKAVLALSGGLDSMFLFKVLSTYQKELEIELILAHVNHKQRIESDWEKELRKLAEEAELPIYISNFS  
GEFSEARARNFRYDFQEVMMKKTGATALTVAHADDQVETIFMRLIRGTRRLYLSGIKEKQVVGEEIIRPFLHFQKKDFPSIFHFED  
TSNQENHYFRNRIRNSYLPELEKENPRFRDAILGIGNEILDYDLAIELSNINVEDLQQLFSYSESTQVLLQTYLNRFPDLNLTKA  
QFAEVQQLKSKSQYRHPKNGYELIKEYQQFQICKISPQADEKEDELVLHYQNQVA YQGYLFSFGLPLEGELIQIPVSRETSIHIRH  
RKTGDVLKNGHRKKLRLFIDLKIPMEKRNSALIEQFGEIVSILGIATNNLSKTKNDIMNTVLYIEKIDRZ

(SEQ. ID. NO. 25)

MRKFLIILLPSFLTISKVVSTEKEVVYTSKEIYYLSQSDFGIYFREKLSSPMVYGEVPVYANEDLVESGKLTPTKTSFQITEWRLNK  
QGIPVFKLSNHQFIAADKRFLYDQSEVTPTIKK VWLESDFKLYNSPYDLKEVKSSLSAYSQVSDIKTMFVEGREFLHIDQAGWVAK  
ESTSEEDNRMSKVQEMLEKYQKDSFSIYVKQLTTGKEAGINQDEKMYAASVLKLSYLYYTQEKINEGLYQLDITTVKYVSAVND  
PGSYKPEGSGSLPKKEDNKEYSLKDLITKVSKESDNVAHNLLGYISNQSDATFKSKMSAIMGDDWDPKEKLISSKMAGKFMEAI  
YNQNGFVLES LTKTDFDSQRIAKGVSVKVAHKIGDADEFKHD TG VYADSPFILSIFTKNSDYDTISKIAKDVEYEV LKZ

(SEQ. ID. NO. 26)

MKKQNNGLIKNPFLWLLFIFFLVTFQYFYSNGNSGGSQQINYTELVQEITDGNVKELTYQPNGSVIEVSGVYKNPKTSKEETGIQF  
FTPSVTKVEKFTSTILPADTTVSELQKLATDHKA ETVKHESSGIWINLLVSIVPFGILFFFLFSMMGMNMGCGNGRNPMSFGRSKA  
KAANKEDIKVRFSVAGAE EEEKQELVEVVEFLKDKPRFTKLGARIPAGVLLGPPGTGKTLAKAVAGEAGVPFFSISGSDFVEMF  
VGVGASRVRS LFEDAKKAAPAIIFIDEIDAVGRQRGVGLGGNDEREQTLNQLLIEMDGFEGNEGHVIAATNRSDVLPDALLRPGR  
FDRKVLVGRPDVKGREAILKVHAKNKPLAEDVDLKLVAQQTGPGFVADLENVLNEALVAARRNKSII DASIDEAEDRVIAGPS  
KKDKTVSQKERELVAYHEAGHTIVGLVLSNARVVHKVTIVPRGRAGGYMIALPKEDQMLLSKEDMKEQLAGLMGGRVAEEIIFN  
VQTTGASNDFEQATQMARAMVTEYGMSEKLGPVQYEGNHAMLAGQSPQKSISEQTAYEIDEEVRSLLNEARNKAAEIIQSNRETH  
KLIAEALLKYETLDSTQIKALYETGKMPEAVEESHALS YDEVKSKMNDKZ

(SEQ. ID. NO. 27)

MKRSSLLVRMVISIFLVFLILLALVGTFYYQSSSAIEATIEGNSQTTISQTSHFQSYIKKLETTSTGLTQQTDLVAY AENPSQDKVEG  
IRDLFTILKSDKDLKTVVLVTKSGQVISTDDSVQMKTSDDMMAEDWYQKAIHQGAMPVLT PARKSDSQWVISVTQELVDAKGA  
NLGVRLRLDISYETLEAYLNQLQLGQGFIFIENHEFVYHPQHTVYSSSSKMEAMKPYIDTGQGYTPGHKSYVSQEKIAGTDWTV  
LGVSSEKLDQVRSQQLWTLGASVTSLLVCLLVWFS LKRWIAPLKDRLRETMLEIASGAQNLR AKEVGAYELREVTRQFNAML D  
QIDQLMVAIRSQEETTRQYQLQALSSQINPHFLYNTLDTIHWMAEFHDSQRVVQVTKSLATYFRLALNQGDICLSDEINHVRQYL

FIQKQRYGDKLEYEINENVAFDNLVLPKLVLPQPLVENALYHGHIKEKEGGHIKLSVQKQDSGLVIRIEDDGVGFQDAGDSSQSQLK  
RGGVGLQNVQDQRLKLHFGANYHMKIDSRPQKGTKVEIYNRIETSZ

## (SEQ. ID. NO. 28)

MKRSSLLVRMVISIFLVFLILLALVGTFFYYQSSSSAIEATIEGNSQTTISQTSHFQSYIKKLETTSTGLTQQTDLVLAENPSQDKVEG  
IRDLFLTLKSDKDLKTVVLTGSGQVISTDDSVQMKTSDDMMAEDWYQKAIHQGAMPVLTTPARKSDSQWVISVTQELVDAKGA  
NLGVLRLDISYETLEAYLNQLQLGQQGFAGFIENNEHFVYHPQHTVYSSSSKMEAMKPYIDTGQGYTPGHKSYSVSQEKIAGTDWTV  
LGVSSLEKLDQVRSQLLWTLGASVTSLLVCLCLVWFSCLKRWIAPLKLRETMLEIASGAQNLRAKEVGAYELREVTRQFNAMLD  
QIDQLMVAIRSQEETTRQYQLQALSSQINPHFLYNTLDTIWMMAEFHDSQRVVQVTKSLATYFRLALNQKDKLCLSDIENHVRQYL  
FIQKQRYGDKLEYEINENVAFDNLVLPKLVLPQPLVENALYHGHIKEKEGGHIKLSVQKQDSGLVIRIEDDGVGFQDAGDSSQSQLK  
RGGVGLQNVQDQRLKLHFGANYHMKIDSRPQKGTKVEIYNRIETSZ

## (SEQ. ID. NO. 29)

MFFKLLREALKVKQVRSKILFTIFIVLVFRIGTSITVPGVNANSLNALSGLSFLNMLSLVSGNALKNFSIFALGVSPYITASIVVQLLQ  
MDILPKFVWEGKQGEVGRRLNQATRYIALVLAQVQSIGITAGFNTLAGAQLIKTALTPQVFLTIGIILTAGSMIVTWLGEQITDKG  
YGNVSMIIFAGIVSSIPEMIQGIYVDYFVNVPSRRITSSIFVILIITVLLIYFTTYVQAEYKPIQYTKVAQAGAPSSSYLPLKVNPA  
VIPVIFASSITAAPAILQFLSATGHDWA WVRVAQEMLATTSPTGIAMYALLIILFTFFYTFVQINPEKAAETYKRVPVISMFEVLVK  
VQKNICLNFFVVLQLLVPSSLVZ

## (SEQ. ID. NO. 30)

MDIRQVTETIAMIEEQNFDIRTITMGISLLDCIDPDINRAAEKIYQKITTKAANLVAVGDEIAAELGIPVKNRVSVPISLIGAATDAT  
DYVVLAKALDKAAKEIGVDFIGGFSALVQKGYQKQDEILINSIPRALAETDKVCSSVNIQSTKSGINMTAVADMGRHIIKETANLSD  
MGVAKLVVFANAVEDNPFMAGAFHGVGEADVIVNVGSGPGVVKRALEKVRGQSFVVAETVKKTAFKITRIGQLVGQMASERL  
GVEFGIVDLSLAPTPAVGDSVARVLEEMGLETVGTHGTTAALALLNDQVKKGGVMACNQVGGLSGAFIPVSEDEGMIAAVQNGS  
LNLEKLEAMTAICSVGLDMIAIPEDTPAETIAAMIADEAAIGVINMKTAVRIIPKKGEGDMIEFGGLGTAPVMKVNAGASSVDFISR  
GGQIPAPIHSFKNZ

## (SEQ. ID. NO. 31)

MTQIIDGKALAAKQQLAEKTAKLKEETGLVPGLVVILVGDNPASQVYVRNKERSALAAGFRSEVVRVPETITQEELLDLIAKYN  
QDPAWHGILVQLPLKHIDEEAVLLAIDPEKDVDFHPLNMGRWLSGHPVMIPTPAGIMEMFHEYIDLECKNAVIVIGRSNIVGK  
PMAQLLLAKNATVTLTSHRTHNLKVAADILVVAIGRAKFVTADFVKPGAVVIDVGMNRDENGKLCGDVDYEA VAPLASHIT  
PVPGGVGPMTITMLMEQTYQAALRTLDRKZ

## (SEQ. ID. NO. 32)

MSKFNRHILVVLDSVGIGAAPDANNFVNAGVDPDGASDTLGHISKTVGLNVPNMAKIGLGNIPRETPLKTVAAESNPTGYATKLEEV  
SLGKDTMTGHWIEMGLNITEPFDTFWNGFPPEILTKEIEFSGRKVIREANKPYSGTAVIYDFGPRQMETGELIITYSADPVLQIAAHE  
DIIPDELRYRICEYARSITLERPALLGRIIARPYVGEFGNFTRTANRRDLAVSPFFPTVLDKLEAGIDTYAVGKINDIFNGAGINHDM  
GHKNSNSHGIDTLLKTMGLAEFEKGFSFTNLVDFDALYGHRRNAHGYRDCLEHFERLPEIIAAMRENDLLITADHGNDPTYAG  
TDHTREYIPLLAYSPAFKGNGLIPVGHFADISATVADNFGVETAMIGESFLDKLVZ

## (SEQ. ID. NO. 33)

MFISISAGIVTFLTLVEIPAFIQFYRKAQITGQQMHEDVKQHQAAGTPTMGGVLFLITSVLVAFFFALFSSQFSNNVGMILFILVLY  
GLVGLDDFLKVFVKINEGLNPKQKLALQLLGGVIFYLYFERGGDILSVFGYPVHLGFFYIFFALFWLVGFSNAVNLTGVDGLASI  
SVVISLSAYGVIAVYVQGMIDILLVILAMIGGLGFFIFNHKPAKVFMDVGSLSALGGLAAISMAHJQEWETLLIIGIVYVFETTSVM  
MQVSFYKLTGGKRIFRMTPVHHHFELGGLSGKGNPWEWKVDFFWVGVLASLLTLAILYLMZ

## (SEQ. ID. NO. 34)

LFKKNKDILNIALPAMGENFLQMLMGMVDSYLAHLGLIAISGSVAGNIITIYQAIFIALGAAISSVISKSIGQKQDQSKLAYHVTEA  
LKITLLLSFLLGFLSIFAGKEMIGLLGTERDVAESGGYLSLVGGSIVLLGLMTSLGALIRATHNPRPLVVSFLSNALNIFSSLAIFV  
LDMGIAGVAVGTIVSRVLVGLVILWSQLKLPYGPPTFGLDKELLTALPAAGERLMMRAGDVVIALVVSFGTEAVAGNAIGEVL  
TFNYMPAFGVATATVMLLARAVGEDDWKRVASLSKQTFWLSLFLMLPLSFSIYVLGVPLTHLYTTDSLAVEASVLVTLFSLGLTP  
MTGTVIYTA VVQGLGNARLPFYATSIGMWCIRIGTGylmGIVLGWGLPGI WAGSLLDN GFRWFLR YRYQRYMSLK GZ

## (SEQ. ID. NO. 35)

MQTQEKHSQAALVGLQHLLAMYSGSILVPIMIALGYSAEQLTYLISTDIFMCGVATFLQLQLNKYFGIGLPPVVLGVAFQSVAPLI  
MIGQSHGSGAMFGALIASGIYVVLVSGIFSKVANLFPISVTGSVITITGLTLPV AIGNMGNNVPEPTGQSLLAAITVLIILLINIFTKG  
FKISISILIGLVGTAAATMGLVDFSPVAVAPLVHVPTPLYFGMPTFEISSIVMMCIATVSMVESTGVYALSDITKDPIDSTRLRNG  
YRAEGLAVLLGGIFNTFPYTGFSQNVGLVKLSGIKKRLPIYYAAGFLVLLGLLPKFGALAQIIPSSVLGGAMLMVFGFVSIQGMQIL  
ARVDFANNEHNFLIAAVSIAAGVGLNNSNLFVSMPTAFQMFFSNGIVVASLLAIVLNAVNLNHHKKKZ

## (SEQ. ID. NO. 36)

MKDRIKEYLQDKGKVTVDNLAQALGKDSSKDFRELIKTSLMERKHQIRFEEDGSLTLEIKKKHEITLKGIFHAHKNNGFGVFSLEGE  
EDDLFVGKNDVNYAIDGDTVEVVIKK VADRNGKGTAAEAKEIIDILEHSLTTVVGVQIVLDQEKPKYAGYIRSKNQKISQPIYVKKPAL  
KLEGTEVLKVFIDKYPKSKHDFVAVSLDVVGHSTDVGDVLEVLSEMDIVSEFPEAVVKEAESVPDAPSQKDMEGRLDLRDEITF  
TIDGADAKDLDDAVHIKALKNGNLEFGVHIADVSYVTEGSALDKEALNRATS VYVTRVVPMLPERLSNGICSLNPQVDRLTQS  
AIMEIDKHGRV VNYTITQTVIKTSFRMTYSDVNDILAGDEEKREYHKIVSSIELMAKLHETLENMRVKGALNFDTPNEAKILVDK  
QGKPVDIVLRQRGIAERMIESFMLMANETVAEHFSKLDLPFIYRIHEEPKAEKVQKFIDYASSFGLRIYGTASEISQEALQDIMRAVE  
GEPYADVLSMMLLRSMQQARYSEHNHGHYGLAADYYTHFTSPIRRYPDLLVHRMIRDYGRSKEIAEHFEQVPEIATQSSNRERRA  
IEAEREVEAMKKAEMYEEYVGEEDYDAVSSIVKFLFVELPNTVEGLIHITNLPEFYHFNERDLTLRGEKSGITFRVGQIRIRVERA  
DKMTGEIDFSFVPEFVDIEKGLKQSSRSRGRDRSNRRSDKKEDKRKSGRSNDKRKHSQKDKKKKGKPKFYKEVAKKGAKHGK  
RGKGRRTKZ

## (SEQ. ID. NO. 37)

MGTGTGTIIDLIIIVYLLAVLVAGIYFSKKEMKGKEFFKGDGSPVWYVTSVSIFATMLSPISFLGLAGSSYAGSWILWFAQLGMVV  
AIPLTIRFILPFIARIDIDTAYDYLDKRFNSKALRIISALLFIYQLGRMSIIMYLPASGLSVLTGIDINILIMGVVAIVYSYTGGKLSVL

WTDFIQGVILISGVVLALFVLIANIKGGFGAVAETLANGKFLAANEKLFDPNLLSNSIFLIVMGSGFTILSSYASSQDLVQRFTTTQNI  
KKLNKMLFTNGVLSLATATVFYLI GTGLYVFFYQVQNADSAASNIPQDQIFMYFIA YQLPVGITGLILAAIY AASQSTISTGLNSVATS  
WTLDIQDVISKNMDSNRRTKIAQFVSLAVGLFSIGVSIVMAHSDIKSAYEWFNSFMGLVLGLLGGVFILGFVSKKANKQGAYAALI  
VSTIVMVFICYFLPPTAVSYWAYSLISISVSVVSGYIVSVLTGNKVSAPKYTTIHDITEIKADSSWEVRHZ

(SEQ. ID. NO. 38)

MKFSKKYIAAGSAVIVSLSCAYALNQHRSQENKDNRRVSVYVDGSSQSKSENLTDPQVSQKEGIAEQIVIKITDQGYVTSHGDH  
YHYNGKVPYDALFSEELLMKDPNYQLKDADIVNEVKGGYIHKVDGKYVYVLKDAHADNVRTKDEINRQKQEHVKDNEKVNS  
NVAVARSQGRYTTNDGYVFNPAHIEDTGNAYIVPHGGHYHYIPKSDLSASELAAAKAHLAGKNMQPSQLSYSSTASDNNTQSV  
KGSTSKPANKSENQLSLLKELYDSPAQRYESDGLVFDPAKISRTPNGVAIPHGDHYHFIPYSKLSALEEKIARMVPISGTGSTVST  
NAKPNEVVSSLGSLSSNPSSLTTSKELSSASDGYIFNPKDIVEETATAYIVRHGDHGFHYIPKSNQIGQPTLPNNSLATPSPSLPINPGTS  
HEKHEEDGYGFDANRIIAEDESFGVMVSHGDHNYFFKKDLTEEQIKVRKNIZ

(SEQ. ID. NO. 39)

MKKRAIVAVIVLLIGLDQLVKSIVYQQIPLGEVRSWIPNFVSLTYLQNRGAAFSILQDQQLFAVITLVVVIGAIWYLHKHMEDSF  
WMVLGLTLIIAGGLGNFIDRVSQGFVDMFHLDIFNFAIFNVADSYLTGVIIIAMIAMLKKEINGNZ

(SEQ. ID. NO. 40)

MNTNLASFIVGLIIDENDRFYVQKDGQTYALAKEEGQHTVGDYVKGFAITDMKQKLRLTTLEVTATQDQFGWGRVTEVRKDLG  
VFVDITGLPDKIVVSLDILPELWPKKGDQLYIRLEVDDKDRIWGLLAYQEDFQRLARPAYNNMQNQNPVPAIVYRLKLSGTFV  
YLPENNMLGFIHPSERYAEPRLGQVLDARVIGFREVDRTLNLSLKPRSFEMLENDQAQMILTYLESNGGFMTLNDKSSPDDIKATFGI  
SKGQFKKALGGLMKAGKIKQDQFGTELIZ

(SEQ. ID. NO. 41)

MKDVSFLKKVFKSRNLNWIVLALFVSVLGVTFYLSQNTANSLSLESRLSRIANERAINENEEKLSQMSDTSSEYQFAKNLD  
VQKNLLTRKTEILTLLKEGRWKEAYYLQWQDEEKNEYFVSNPDPTASPLKMGVDRERKIYQALYPLNIKAHTLEFPTHGIDQIVWI  
LEVIPSLFVVAIIFMLTQLFAERYQNHLDTAHLYPVSKVTFAISSLGVGVGYYTVLFIGICGFSFLVGSLSISGFGQLDYPYPIYSLVNQ  
EVTIGIKIQDVLFPGLLLAFLAFIVIVEVYLYIAFFKQKMPVFLSLIGIVGLLFGIQTIQPLQRIAHLPFTYLRVSEILSGRLPKQIDNV  
DLNWSMGMVLLPCLIIIFLLGILFIERWGSSQKKEFFNRFZ

(SEQ. ID. NO. 42)

MMKFILDIVSTPAILVALIAILGLVLQKKLPDIKGGIKTFVGFLVSVGGAGIVQNSLNPFGTMFEHAFHLSGVVPNNEAIVAVALT  
TYGSATAMIMFAGMVFNILIRFTRFKYIFLTGHHTLYMACMIAVILSVAGFTSLPLILLGGLALGIMSISPAFVQKYMVQLTGNDK  
VALGHFSSLGYWLSGFTGSLIGDKSKSTEDIKFKSLAFLRDSTVSITLSMAVYIIVAIFAGSEYIEKEISSGTSGLVYALQLAGQFAA  
GVFVILAGVRLILGEIVPAFKGISERLVPNSKPAIDCPIVYTYAPNAVILGFISSFVGGVLSMVIMIASGTVVILPGVVPHFCCGATAG  
VIGNASGGVRGATIGAFQLGILISFLPVFLMPVLGGLGFGQSTFSDADFLGSGILGMLNQFGSQAGIVIGLVILAVMFGVSFIKKPS  
ATEEZ

(SEQ. ID. NO. 43)

MIKTFLSALSIVLFSIPIITYSFFSSNLNIWLSTQPILAQIYAFPLATATMAAAILSFLFFLSFYKKNKQIRFYSGILLLSLILLFGTDK  
TSSASNKTITLKLVTWNVANQIEAQHIERIFSHFDADMAIFPELATNIRGEQENQRIKLLFHQVGLSMANYDIFTSPTNSGIAPVT  
VIVKKS YGFYTEAKTFHTTRFGTIVLHRSRKNIPDIIALHTAPPLPGLMEIWQKDLNIIHNQLASKYPKAIAGDFNATMRHGALAKI  
SSHRDALNALPPFERGTWNSQSPKLFNATIDHILLPKNHYYVKDLDIVSFQNSDHRCIFTEITFZ

(SEQ. ID. NO. 44)

MNPIQRSWAYVSRKRLRSFILFLILLVLLAGISACLTLMKSNKTVESNLYKSLNTSFSIKKIENGQTFKLSDLASVSKIKGLENVSP  
ETVAKLKDKAEVTEGEQSVERRDLSAADNNLVSLTALEDSSKDVFTTSSAFNLKEGRHLQKGDSSKILIHHEELAKKNGLSLHDKIGL  
DAGQSESGKQGTVEFIIGFSGKKQEKFTGLSSDSENQVFTDYESSQTLLGNSEAQVSAARFYVENPKEMDGLMKQVENLAL  
QGYQVEKENKA FEQIKDSVATFQTFLTIFLYGMLIAGAGALILVLSLWLRERVYEVGILLALGKGGSSIFLQFCLEVVVLVSLGALLP  
AFVAGNAITTYLLQTLASGDQASLQDTLAKASSLTSILSFAESYVFLVLLSCLSVLALCFLFLFRKSPKEILSSISZ

(SEQ. ID. NO. 45)

MLHNFAFYVTRKFFKSIVIFLIILLMASLSLVGLSIKGATAKASQETFKNITNSFSMQINRRVNQGTGPRGAGNIKGEDIKKIKTENKAIE  
SYVKRINAIGDLTGIDYDIETPETKKNLTADRKRFGSSLMITGVNDSSKEDKFVSGSYKLVEGEHLTNDKDKILLHKDLAAKHG  
WKVGDKVKLDSNIYDADNEKGAKETVEVTIKGLFDGHNSAVTYSQELYENTAITDIHTAAKLYGYTEDTAIYGDATFFVTADK  
NLDDVMKELNGISGINWKS YTLVKSSSNYPALQESISGMYKMANLLFWGSLSFVLLALLLSLWINARRKEVGILLSIGLKQASIL  
GQFITESILIAIPALVSAYFLANYTARAIGNTVLANVTSGVAKQASKAAQASNLGGGAEVDFGSKTLLSSLDISIQTSDFIIFVLALVLV  
VLVMALASSNLLRKQPKELLDGEZ

(SEQ. ID. NO. 46)

MSQDKQMKAVPSLLQRVINISSIVGGVGLSIFCIWAYQAGILQSKETLSAFIQQAGIWGPPLFIFLQILQTVVPIPGALTSVAGVFIYG  
HIIGTIYNYIGIVIGCAIIFYLVRLYGAAAFVQSVVSKRTYDKYIDWLDKGNRFRDRFFIFMMIWPISPADFLCMLAALT KMSFKRYMTII  
ILTKPFTLVVYTYGLTYIIDFFWQMLZ

(SEQ. ID. NO. 47)

MRNMVWVVIKETYLRHVESWSFFFVVISPFLLGISVGIGHLQGSSMAKNNKVAVVTTVPVSAEGLKNVNGVNFYKDEASAKEAI  
KEEKLKGYLTIDQEDSVLKAVYHGETSLENGIKFEVTGTLNELQNQLNRSTASLSQEQEKRLAQTIQFTEKIDEAKENKKFIQTIAA  
GALGFFLYMILITYAGVTAQEVASEKGTKIMEVVFSSIRASHYFYARMMAFLFVILTHIGIYVVGGLAAVLLFKDLPFLAQSGILDH  
LGDALSINTLLFILISFLVYVLAFLGSMVSRPDSKQALSPMLIMLGGFFGTALGAAGDNLKIGSYIPFISTFFMPFRFTINDY  
AGGAEAWISLAITVIFAVVATGFIGRMYASLVLTQDLDGIWKTFRKALSYKZ

(SEQ. ID. NO. 48)

MTETIKLMKAHTSVRRFKEQEIPQVDLNEILTAAQMASSWKNFQSYSVIVVRSQEKKDALYELVPQEAIRQSAVFLLVGDLNRAE  
KGARLHTDTFQPQGVGELLISSVDAALAGQNALLAAESLGYGGVIGLVRYKSEEV AELFNLPDYTYSVFGMALGVPNQHHDMKP

RLPLENVVFEEYQEOSTEAIQAYDRVQADYAGARATTSWSORLAEQFGQAEPSSTRKNLEQKKLLZMLKLIIVGTNSKRSTNR  
QLLQYMQKHFTDKAEIELVEIKAIPVFNKPADKQVPAEILEIAAKIEEADGVIIIGTPEYDHSIPAVLMSALAWLSYGIYPLLNKPMIT  
GASYGTLGSSRAQLQLRQILNAPEIKANVLPDEFLLSHSLQAFNPSPGDLVDLDVIKKLDAIFDDFRIFVKITEKL RNAQELLRKDAED  
FDWENZ

(SEQ. ID. NO. 49)

MNTYQLNNGVEIPVLGFGTFKAKDGEEAYRAVLEALKAGYRHIDTAAYQNEESVGQAIKDSGVPREEMFVTTKLWNSQQTYEQ  
TRQALEKSIEKLGLDYLILYHWPNPPLRENDWKTNAEVRAMEDLYQEGKIRAIGVSNFLPHHLDALLETATIVPAVNQV  
RLAPGVYQDQVVAYCREKGILLEAWGPFQGGELFDSKQVQEIAANHGKSVAQIALAWSLAEGFLPLPKSVTTSRIQANLDCFGIEL  
SHEERETLKTIQVQSGAPRVDDVDFZ

(SEQ. ID. NO. 50)

MRCKMLDPIAIQGLPAIRWYALCIVTGLILAVYLTMKEAPRKKIIPDDILDFILVAFPLAILGARLYYVIFRFDYYSQNLGEIFAIWN  
GGLAIYGGITGALVLYIFADRKLINTWDFLDIAAPSVMAIQLSGRWGNFFNQEAYGATVDNLDYLPGFIRDQMYIEGSYRQPTFL  
YESLWNLGFGALILIFRRKWKSLRRGHITAFYLIWYGFGRMVIIEGMRDLSLFFGFRVSQWLSVVLIGLGIMIVYQNRKKAPYYIT  
EENZ

(SEQ. ID. NO. 51)

MGKLSSILLGTVSGAALALFLTSDKGKQVCSQAQDFLDDLREDPEYAKEQVCEKLTVEVKEQATDFVLKTKEQVESGEITVDSILAQ  
TKSYAFQATEASKNQLNNLKEQWQEKAEALDDSEEIVIDITEEZ

(SEQ. ID. NO. 52)

MKTKLIFWGSMLFLLSILLSITTYLAWIFYPMIEIQLWNLNTRVYLKPETIQYNFHLNMNYLTNPFSQVQLQMPDFRSSAAGLHHFAV  
KNLHFLVQLVALVTLPSFYVFNRIKDFLSLYRKSLLALVVLPMIGLGGVLIGFDQFFTLFHQILFVGGDTWLFDPKADPVI  
LPETFFLHAFLLFFALYENFFGYLYLKSRRKZ

(SEQ. ID. NO. 53)

MTYHFTTEYDIIVIGAGHAGVEASLAASRMGCKVLLATINIEMLAFMPCNPISIGGSAGKIVVREVDALGGEMAKTIDKTYIQMKML  
NTGKGPAVRALRAQADKELYSKEMRKTVENQENLTLRQTMIDEILVEDGKVVGVRTATHQEYAAKAVIVTTGTALRGIIIGDLK  
YSSGPNHSLASINLADNLKELGLEIGRFKTGTPPRVKASSINYDVTEIQPGDEVPHFSYTSRDEYVKDQVPCWLTNTNGTSHEIIQ  
NNLHRAPMFTGVVKGVPYCPYPSIEDKIVRFADKERHQLFLEPEGRNTEEVYVQGLSTSLPEDVQRDLVHSIKGLENAEMMRGT  
AIEYDMVLPQLRATLETKKISGLFTAGQTNGTSGYEEAAGQGIIAGINAALKIQGKPELILKRSYGIVGMIDDLVTKGTIEPYRL  
TSRAEYRLILRHDNADMRLTEMGREIGLVDDERWARFEIKKNQFDNEMKRLDSIKLKPVKETNAKVEEMGFKPLTDAVTAKEFLR  
RPEVSYQDVVAFIGPAAEDLDDKIIELIETIEIKYEGYISKAMDQVAKMKRMEEKRIPANIDWDDIDSIAETEARQKFKLINPETIGQAS  
RISGVNPADISILMVYLEGKNRSISKTLQKSKZ

(SEQ. ID. NO. 54)

MTKQVLLVDDDEHILKLLDYHLSKEGFSTQLVTNGRKALALAEETPFDFILLDMLPQLDGMVECKRLRAKGVKTPIMMVSAKSD  
EFDKVLALLEGADDYLTKPFSPRELLARVKAFLRRTKGEQEGDDSDNIADDSWLFGLTKVYPERHEVYKANKLLSLTPKEFESDK  
NPFEVFKVSKVTAQZ

(SEQ. ID. NO. 55)

MTTFKDGFLWGGAVAAHQLEGGWQEGGKGISVADVMTAGRHGVAREITLGVLEGKYYPNHEAIDFYHRYKEDIALFAEMGFKC  
FRTSIAWTRIFPKGDELEPNNEGLQFYDNLFDECLKNGIEPVITLSHFEMPYHLVTEYGGWKNRKLIDFFARFAEVVFKRYKDKVK  
YWMFTNEINNQANYQEDFAPFTNSGIVYEEGDNREAIMYQAAHYELVASARAVKIGHEINPDFQIGCMIAMCPIYPVTCNPKDILM  
AMKAMQKRYFADVHVLGKYPEHIFKYWERKGISVDFTAQDKEDLLGGTVDYIGFSYMSFAIDSHRENNPYFDYLETEDLVKN  
NYVKASEWEWQIDPEGLRYALNWFTDHYHLPLFIVENGFGAIDQVAADGMVHDDYRIEYLGAIHIREMKKAVVEDGVDLMGYTP  
WGCIDLVSAGTGEMRKRYGFIYVDKDDNGKGSYNRSPKKSFGWYKEVISSNGESVEZ

(SEQ. ID. NO. 56)

MDQQNGLFGFLENHVMGPMGKLAQFKVVRITAAGMAAVPFTIVGSMFLVFSILPQAFSFWPIVADIFSASFDFKFTSLYMVANYA  
TMGSLSLYFVLSLAYELTKIYAEEEEELNMNPLNGALLALMAFVMTVPQIIFDGGMMKTVTSLKEGAVIADGWAMGNVVARFGTT  
GIFTAIIAIVTVLIYRMCVKNHWIKMPEAVPEGVSRGFTALVPGFVAVFVIFINGLLVAMGTDIFKVIAPFGFVSNLTSNWIGL  
MIYLLTQLLWIVGIHGANIVFAVSPIALANMAENAAGGHFAVAGEFSNMFVIAGGSGATLGLCLYIAFASKSEQLKAIGRASVVP  
ALFNINEPLIFGLPIIYNPALAIPFILAPMVTATIIYVANSNLFKPIIAQVWPPTVGIGAFLGTADLRAVLVALVCAFAFLVYLPFIR  
VYDQKLVKEEQGIZ

(SEQ. ID. NO. 57)

MKKFYVSPPIFPLVGLIAFGVLSTFIIFVNNNLLTVLILFLFVGGYVFLFKKL RVHYTRSDVEQIYVNHQAEESLTALLEQMPVGVM  
KLNLSGGEVWFNPAELILTKEDGDFLEAVQTIKASVGNPSTYAKLGEKRYAVHMDASSGVLYFVDVSREQAITDELVTSRPV  
IGIVSDNYDDLEDETSEDISQINSFVANSIFSEKHMMSRRVSMDFYLFDTYTVLEGLMNDKFSVIDAFREESKQRQLPLTSL  
MGFSYDGNHDEIGKALLNLNLAEVRGGDQVVVKENDETKNPVYFGGSAASIKRTRTRTRAMMTAISDKIRSVQVVFVGHK  
NLDMDALGSVGMQLFASNVIENSYALYDEEQMSPDIERAVSFIEKEGVTKLLSVKDAMGMVTNRSLLILVDHSKALTLSKEFY  
DLFTQTVIDHHRDQDFPDNAVITYIESGASSASELVTELIQFQNSKKNRSLRMQASVLMAGMMLDTKNFTSRVTSRTFDVASYL  
RTRGSDSIAIQEIAATDFEYREVNELILQGRKLGSVDVLAIEAKDMCYDTVVISKAADAMLAMSGIEASFVLAKNTQGFISISARS  
SKLNVQRIMEELGGGGHFNLAQAQIKDVTLSEAGEKLTIVLNMKEKEKEEZ

(SEQ. ID. NO. 58)

MKEKNMWKELLNRRAGWILVFLAVLLYQVPLVVTSLTLKEVALLQSGLIVAGLSIVVLALFIMGARKTKLASFNFSFFRAKDLAR  
LGLSYLVIVGNSILQSILLQLSNETTTANQSQINDMVQNSSLISFFLLALLAPICEILCRGIVPKKIFRGKENLGFVVGTVFALLHQP  
SNLPSLLIYGGMSTVLSWTAYKTQRLEMSILLHMIVNGIAFCLLALVIMSRTLGISVZ

(SEQ. ID. NO. 59)

MKEKNMWKELLNRAWILVFLAVLLYQVPLVVTSTILTKEVALLQSGSLIVAGLSIVVLALFIMGARKTKLASFNFSFFRAKDLAR  
LGLSYLVIVGNSILGSIILLQSNETTANQSQINDMVQNSSLSFFLLALLAPICEEILCRGIVPKIFRGKENLGFVVGTVFALLHQF  
SNLPSLLIYGGMSTVLSWTAYKTQRLEMSILLHMIVNGIAFCLLALVIMSRTLGISVZ

(SEQ. ID. NO. 60)

MDTQKIEAAVKMIEAVGEDANREGLQETPARVARMYQEIFSGLGQTAEHLSKSFEIIDDNMVVEKDIFFHTMCEHHFLPFYGRA  
HIAYPDGRVAGLSKLARTVEVYSKKPQIQERLNIEVADALMDYLGAKGAFVVEIAEHMCMMSMRGVKPGTATLTTVARGLFETD  
KDLRDQAYRLMGLZMKDLFLKRKQAFRKECLGYLRVYVNDHFVFLFLLVLLGFLAYQYSQLLQHFPENHWIPILLFVGITSVLLLLW  
GGTATYMEAPDKLFLLVGEEIEKLHLKRQTGISLVFWLFLVQTLFLLFAPLFAMGYGLPVFLLYVLLGVGKYFHFQKASKFT  
ETGLDWDYVISQESKRKQVLLRFFALFTQVKGISNSVKRRAYLDFILKAVQKVPKGIWQNLVLRSLRNGDLFALSRLRLLSLLA  
QVFIEQAWIATAVVVLFNYLLLFQLLALYHAFDYQYLTQLFPLDKGQKEKGLQEVVRGLTSFVLLVELLVGLITFQEKALLALLG  
AGLVLLVLYLPYQVKRQMQDZ

(SEQ. ID. NO. 61)

MRKSIVLAADNAYLIPLLETTIKSVLYHNRDVFYILNSDIAPEWFKLLGRKMEVNVNSTIRSVHIDKELFESYKTGPHINYASYFRFFA  
TEVVESDRVLYLDSIIVTGELATLFEIDLKGYSIGAVDDVYAYEGRKSGFNTGMLLMDVAKWKEHSIVNSLLELAAEQNQVVHL  
GDQSILNIYFEDNWLALDKTYNYMVGIDYHLAQECERLDDNPPTIVHYASHDKPWNTYSSIRLRELWWVYRDLDWSEIAFQRSD  
LNYFERSNQSKKQVMLVTWSADIKHLEYLVQRLPDWHFHLAAPCDCSEELTSLSQYTNVTYQNVLHRSRIDWLLDDSIYVLDINT  
GGEVFNVTTRAQESGKKIFAFDITRKSMDGGLYDGIFSVERPDDLVDKMKNIEIEZ

(SEQ. ID. NO. 62)

MTKIYSSIAVKKGLFTSFLFIYVLGSRILPFVDLNTKDFLGGSTAYLAFSAALTGGNLRSLSIFSVGLSPWMSAMILWQMFSFSKRL  
GLTSTSIEIQDRRKMYLTLLIAVIQSLAVSLRPLVQSSYSAILVVLMTILLIAGTFFLVWLSDLNASMGIGGSIVILLSSMVLNIPQDV  
LETFQTVHIPTGHIIVLLALLTLVFSYLLALMYRARYLVPVNKIGLHNRFRKYSYLEIMLNPAGGMPYMYVMSFLSVPAYLFILLGFIF  
PNHSLAALSKEFMVGKPLWVYVYISVLFSLIIFAFVTMNGEEIADRMKKSGEYIYGIYPGADTSRFINRLVLRFSVIGGLFNVIMA  
GGPMLFVLFDEKLLRLAMIPGLFMMFGGMIFTIRDEVKALRLNETYRPLIZ

(SEQ. ID. NO. 63)

MSSLSQDELVAKTVEFRQRLSEGESLDDILVEAFVREADKRILGMFPYDVQVMGAIVMHYGNVAEMNTGEGKTLTATMPVYL  
NAPSSEGVMVVTNPNEYSKRDAEEMGQVYRFLGLTIGVPFTEDPKKEMKAEKKLIYASDIYTNSNLGFDYLDNDLASNEEGKF  
LRPFNYVIIDEIDILLDSAQETPLIAGSPRVQSNYYAIDTLVTTLVEGEDYIFKEEKEEVWLTTKGAKSAENFLGIDNLYKEEHASF  
ARHLVYAIRAHKLFTKDKDYIIRGNEMVLVDKGTGRLMEMTKLQGGHLQAIEAKEHVKLSPETRAMASITYQSLFKMFNKISGMT  
GTGKVAEKEFIETYNMSVVRIPTNRPRQRIDYPDNLYITLPEKVYASLEYIKQYHAKGNPLLVFVGSVEMSQLYSSLLFREGIAHNV  
LNANNAAREAAQIISSESGQMGAVTVATSMAGRGTDIKLGKGAELGGLIVIGTERMESQRIDLQIRGRSGRQGDPMGSKFFVSLEDD  
VIKKFGPSWVHKKYKDYQVQDMTQPEVLKGRKYRKLVEKAQHASDSAGRSARRQTLEYAESMNIQRDIVYKERNRLIDGSRDLE  
DVVVDIERYTEEVAADHYASRELLFHFIVTNISFHVKEVPDYIDVTDKTAVRSFMKQVIDKELSEKELLNQHDLYEQFLRLSLK  
AIDDNWVEQVDYLQQLSMAIGGQSASQKNPIVEYYQEAAYAGFEAMKEQIHADMVRNLLMGLVEVTPKGEIVTHFPZ

(SEQ. ID. NO. 64)

MIGTFAAALVAVLANFIVPIETPNSANTEIAPPDGIGQVLSNLLKLVDNPNVALLTANYIRILSWAVIFGIAMREASKNSQELLKTI  
ADVTSKIVEWIINLAPFGLGVFKTISDKGVGSLANYGILLVLLVTTMLFVAPVNVNPLIAFFFMRRNPYPLVWNCRLRVSGVTAFFTR  
SSATNIPVNMKLCHDLGLNPDYTSVSIPLGSTINMAGVAITINLLTAAVNTLGPVDFATAFVLSVVAIISCDASGIAGGSLLLIPV  
ACSLFGISNDIAIQIVGVGFVIGVIQDSCETALNSSTDVLFATAVEYAATRKKZ

(SEQ. ID. NO. 65)

MSISQRTTKLILATCLLAYFLNLSSAVSAGIALLSLDTRRSTLKLARNRLFSMLLALAIGVLAFLHLSGFHIWSLGLYLAFYVPL  
AYKMGWEIGTPTSVLVSHLLVQUESTPDLLVNEFLFAIGTGFAALLVNLMPYSREEEIQHYHTLVEEKLKDILQRFKYLSRGDGR  
NRAQLVAELDTLLKEALRLVYLDHSDHLFHQTDYHIHYFEMRQRQSRILRNMAQQINTCHLAASESLILAQLFSKIAQGLSQTNPA  
SDLLDEIERYLEVFRNRSPLKTRFEFETRATLLQLLREAKTFIQVKVDFYQKYRQZ

(SEQ. ID. NO. 66)

MEIMSLAIAVFAVIIGLVIGYVSISAKMKSSQEAELMLLNAEQEATNLRGQAEREADLLVNEAKRESKSLKKEALLEAKEEARKY  
REEVDAEFKSERQELKQIESRLTERATSLDRKDDNLTKEQTLQEQESISDRAKNLDAREEQLEEVEERQKEAELERIGALSQAEAR  
DIIAQTEENLTREIASRIEAEQEVKERSDKMAKDILVQAMQRIAGEYVAESTNSTVHLPPDDTMKGRIIGREGNRNRTFESLTGVDV  
IIDDTPVVTLSGFDPIRREIARMTEMEMLLKDGRIPARIEELVEKNRQEIENKIREYGEAAAYEIGAPNLHPDLMKIMGRQLQFRTSY  
GQNVLRHSIEVAKLAGIMASELGENAALARRAGFLHDIGKAIIDHEVEGSHVEIGMELARKYKEPPVVVNTIASHHGDVEAESVIAV  
IVAAADALSAAARPGARSESLESYIKRLHDLLEEIANGFEGVQTSFALQAGREIRIMVNPGKIKDDKVTLAHKVRKKIENNLDYPGNI  
KVTVIRELRAVDYAKZ

(SEQ. ID. NO. 67)

MMLKPSIDTLLDKVPSKYSVLVILEAKRAHELEAGAPATQGFKSEKSTLRALEEIESGNVTIHPDPEGKREAVRRRIEEERKRKEEEEK  
KIKEQIAKEKEDGEKIZ

(SEQ. ID. NO. 68)

MSAYQLPTVWQDEASNQGAFTGLNRPTAGARFEQNLPKGEQAFQLYSLGTPNGVKVTILLEELLEAGFKEAAYDLYKIAIMDGDQ  
FGSDFVKNLNPNSKIPALLDQSGTENVRVFESAHILLYLAEKFGAFLSPNPVEKVEVLNWLFWQAGAAPFLGGGFGHFFNYAPEKLE  
YPINRFTMEVKRQLDLLDKELAQKPYIAGNDYTIADIAIWSWYGQLVQGNLYQGSAKFLDASSYQNLVKWAEKIANRPAVKRGL  
EVTYTEIKZ

(SEQ. ID. NO. 69)

LASLITSIMFYVGFVDLRDTIQKLSREETVIDPLGATLGHISAAIMFVYLYNTRLSKKSNSNALKAAAKDNLSDAVTSLGTAIAILA  
SSFNYPIVDKLVAIITFFILKTAIDIFISSFSLSGDFDRLLLEDYQKAIMEIPKISKVKSQRGRTYGSNIYLDITLEMNPDLVSFESHEI  
ADQVESMLEERFGVFDTDVHIEPAPIEDEILDNVIYKLLLMREQLIDQGNQLEELLTDFFVYIRQDGEQMDKEAYKTKKELNSAIK  
DIQITSISQKTKLCICYELDGIHITSIWRRHETWQNIHFQETKKEZ



(SEQ. ID. NO. 70)

MTIKLVATMDMGTFLDGNGRFDMDRLKSLLSYKEKGIYFAVASGRGFLSLEKLFAGVRDDIIFIAENGLVEYQGQDLYEATMS  
RDFYLATFEKLKTSPPYVDINKLLLTGKKGSYVLDTVDETYLKVSQHYNENIQKVASLEDITDDIFKFTTNFTEETLEDGEAWVNEN  
VPGVKAMTTGFESIDIVLDYVDKGVAIVELVKKLGITMDQVMAFGDNLNDLHMMQVVGHVPAPENARPEILELAKTVIGHHKKR  
SVIAYMEGLZ

(SEQ. ID. NO. 71)

MADIKLIAIDLDTLLTDDKRLTDRTKETLQAARDRGIKVVLTTGRPLKAMDDFLHELGTGQDEYETITFNGGLVQKNTGEILDK  
TVFSYDDVARLYEETEKLSPDAISEGTVYQIQSDQESLYAKFNPALTFVPVDFEDLSSQMTYNKCVTAFAQEPLDAAIQKISPELF  
DQYEIFKSREMLLEWSPKNVHKATGLAKLISHLGIDQSQVMACGDEANDLSMIEWAGLGVAMQNAVPEVKAAANVTPMTNDE  
EAVAWAIEEYVLKENZ

(SEQ. ID. NO. 72)

MESLLILLLIANLAGFLIWQRQDRQEKHLSKSLEDQADHLSDQLDYRFQARQASQLDQKDLEVVSDDLQEVRIELHQGLTQV  
RQEMTDNLLQTRDKTDQRLQALQESNEQRLEQMRQTVVEEKLEKTLQTRLQASFETVSKQLESVNRGLGEMQTVARDVGALNKV  
LSGKTRKRGILGELQLGQIIEDIMTPAQYEREYATVENSSERVEYAIAKLPQGQDQYVYLPIDSKFPLADYYRLEEAYETGDKDEIERC  
RKSLLASVKRFARDIRNKYIAPPRTTNFGVLFVPTTEGLYSEIVRNPVFFDDLREEQIIVAGPSTLSALLNSLSVGFKTLNIQKSADHIS  
KTLASVKTEFGKFGGILVKAQKHLQHASGNIDELLNRRITAIERTLRHIELSEGEPAIDLLHFQENEEYEDZ

(SEQ. ID. NO. 73)

MKISHMKKDELFEFGFYLIKSADLRQTRAGKNYLAFTFQDDSGEIDGKLWDAQPHNIEAFTAGKVVMKGRREVYNNTQVNOIT  
LRLPQAGEPNPADFKVKSPPVDVKEIRDYMSQMIFKIENPVWQIVRNLYTKYDKEFYSYPAAKTNHHAFETGLAYHTATMVRL  
ADAISEVYPQLNKSLLYAGIMLHDLAKVIELTGPQDQTEYTVRGNLLGHIALIDSEITKTMELGIDDTKEEVLLRHVILSHHGLLE  
YGPSPVRPRIMEAIIHMIDNLDASMMMMSTALALVDKGEMTNKIFAMDNRSFYKPDLDZ

(SEQ. ID. NO. 74)

MSEKAKKGFKMPSSYTVLLIIHIAIMAVLTWFIAPAGAFIEGIYETQPQNPGIWDVLMAPIRAMLGTHPEEGSLIKETSAAIDVAFFILM  
VGGFLGIVNKTGALDVGIAIVKYYKGREKMLILVLMPLFALGGTTYGMGEETMAFYPLLVPVMMAVGFDLSLTGVAIILLGSQIGC  
LASTLNPATGIASATAGVTGDGIVLRLIFWVTLTALSTWVYRYADKIQKDPTKSLVYSTRKEDLKHFNVEESSVESTLSSKQK  
SVLFLFVLFILMVLSPFIPWTDLGVTFDDFNTWLTGLPVIGNIVGSSTALGTWYFPEGAMLFAMGILIGVIYGLKEDKIISSFMNG  
AADLLSVALIVAIARGIQVIMNDGMITDITLWGWKEGLSGLSSQVFIWVYIFYLPMSPFLIPSSSGLASATMGIMAPLGEFVNVPRSLII  
TAYQSASGVNLNLIAPTSGIVMGALALGRINIGTWKFMGKLVVAIIVVTIALLLLTGTFPLZ

(SEQ. ID. NO. 75)

MSNSFVKLLVSQFLANLADIFFRVTHIANIYIISKSVIATSLVPILIGISSFVASLLVPLVTKRLALNRVLSLSQFGKTILLAILVGMFTV  
MQSVAPLVTYLFFVVAISILDGFAAPVSYAIVPRYATDLGKANSALSMTGEAVQLIGWGLGGLLFATIGLLPTTCINLVLYIISFLML  
FLPNAEVEVLESETNLEILLKGWKLVARNPRLRLFVSANLLEIFSNITWVSSIILVFVTELLNKTESYWGYSNTAYSIGIIISGLIAFRLS  
EKFLAAKWEQQLFTPNLKTIONPCLSLDPGWFLFSPNGCFLLDKKEFPYLGISVEKNTKRKETHMNSLPNHHFQNKSFYQLSFDGG  
HLTQYGGILFFQELFSQKLKERISKYLVTDNRYYCRYSDSDILVQFLQLLTGYGTDYACKELSADAYFPKLEGGQLASQPTLS  
RFLSRDEETVHSLRCLNLELVEFFLQFHQLNQLIVDIDSTHFTTYGKQEGVAYNAHYRAHGYHPLYAFEGKTGYCFNAQLRPGN  
RYCSEEDSFTIPVLERFNQLLFRMDSGFATPKLYDLIEKTGQYYLIKLLKNTVLSRLGDLSLPCQDEDLTILPHSAYSETLYQAGS  
WSHKRRVCQFSERKEGNLFYDVISLVNTMTSGTSQDQFQLYRGRGQAENFIKEMKEGFFGDKTDSSTLIKNEVRMMMSCIAYNLY  
LFLKHLAGGDFQTLTIKFRHLFLHVVGKCVTRGRKQLLKLSSLYAYSELSALYSRIRKVNLLNPVPYEPPIRRKASLMMHZ

(SEQ. ID. NO. 76)

MMEFFQQLPHLEPYGNPQYFVYVIAATLPFIQGLFFKKRFAWYEVVLVSFFIVTMLVGGKTNQLAALGIYLCWEILLLLFYKHRYKS  
KDGKWWFYLVSFLSLLPIIFVKVQPAINGTQSLGLGLISYLTFRSVGIVIELRDGVIKDFTLWEFLRFLFMPTFSSGPIDRFKRFNEN  
YQAIPERDELMDMLDESRYIMWGFYKFIHAHVLETLLPPLKNLALQSGGFNLYALAVMYTFGLELFFDFAGYSMPALAIISNL  
MGIRSPINFNKPFLSRDLKEFWNRWHMSLSFWFRDFVFMVMVLTTRKKVFKNRNVTSMAIYVNMIMGFHWGVTWYYIAYG  
LFHGLGLVINDAWVRKKKTLNKKERKKAGKAALPENRWIQLLGMVVTFFHVMLSFLIFSGFLNLLWFKKZ

(SEQ. ID. NO. 77)

MLKRLWMIFGPVLIAGLLVFLLIFFYPTMHNNLGAEKRSVAVATTIDSFKERSQKVRALSDPNVRFVPPFGSSEWLRFDAHPAVL  
AEKYNRSYRYPYLLGQGAASLNQYFGMQMLPQLENKQVYVVISPWFSKNGYDPAAFQYFNGDQLTSFLKHQSGDQASQYA  
ATRLQQFPNVAMKDLVQKLASKEELSTADNEMIELLARFNERQASFFGQFSVRGYVNYDKHVAKYLKILPDQFSYQAIEDVVKA  
DAEKNTSNEMGMENYFYNEQIKKDLKKLSDSKSFTYLSPEYNDLQVLVTQFSKSKVNPFIIPPVNKKWMNYAGLREDMYQ  
QTVQKIRYQLESQGFNTIADFSKDGGEPPFMKDTIHLGWLGWLAFDKAVDPFLSNPTPAPTYHLNERFFSKDWATYDGDVKEFQZ

(SEQ. ID. NO. 78)

MEKNLKALKQTTDQEGPAIEPEKAEDTKTVQNGYFEDAADVDRTLSDYAGNWQSVYPFLEDGTFDQVFDYKAKLTGKMTQAEY  
KAYYTKGYHTDVTKINITDNTMEFVQGGQSKKYTYKYVGKKILTYKKGNRGVRFLFEATDADAGQFKYVQFSDHNVAPVKAEH  
FHFFGGTSQEALFEEDMNWPTYYPDNLSGQIEAQEMLAHZ

(SEQ. ID. NO. 79)

MKDGHLAHHRLLNGRIFQKLLSQDPEALYRGEQKILAVLWNSSETGCATATDIALATGLANNTLTMMIKKLEEQKLIVVSPCGK  
DKRKKYLVLTGKSKQKEVGHVRSQKLDITFYKGFSEEEIHQFEGFQERILANLKEKGNEVZ

(SEQ. ID. NO. 80)

MTNLIATFQDRFSDWLTALSQHLQLSLLTLLAILLAIPLAVFLRYHEKLDWVLQIAGIFQTIPLSALLGLFIPLMGITLPALTALV  
IYAIFPILQNTITGLKGIDPNLQEGAGAFGMTRWERLKKFEIPLAMPVIMSGIRTAAVLIIGTATLAALIGAGGLGSFILLGIDRNNASLI  
LIGALSSAVLAIANFLLVKMEKAKLRTIFSGFALVALLGLSYSPALLVQKEKENLVIAGKIGPEPEILANMYKLLIEENTSMATV  
KPNFGKTSFLYEALKKGIDIDYEFPTGTVTESLLQPSPKVSHEPEQVYQVARDGIAKQDHLAYLKPMYSQNTYAVAVPKKIAQYEG  
LKTISDLKKVEGQLKAGFTLEFNDREDGNKGLQSMYGLNLSNVATIEPALRYQAIQSGDIQITDAYSTDAELERYDLQVLEDDKQLF  
PPYQAGAPLMKEALLKHPELERVLNLTLAGKITESQMSQNLNYQVGVEGKSAKQVAKFEFLQEQLLKKZ

(SEQ. ID. NO. 81)

MMHTYLQKKIENIKTTLGEMSGGYRRMVAAMADLGFSGTMKAIWDDLFAHRSFAQWIYLLVLGSFPLWLELVYEHRIVDWIGMI  
CSLTGIIICVIFVSEGRASNYLFLGINSVIYLLALQKGFYGEVLTTLTYFTVMQPIGLLVWIYQAQFKKEKQEFVARKLDGKGWTKYL  
SISVLWWLAFGFYQSIGANRPYRDSITDATNGVGQILMTAVYREQWIFWAATNVFSIYLWWGESLQIQGKYLILYINSLVGWYQ  
WSKAAKQNTDLLNZ

(SEQ. ID. NO. 82)

MRNMKAKYAVVWAFFLNLTYAIVEFIAGGVFGSSAVLADSVHDLGDAIAIGISAFLETISNREEDNQYTLGKYRFSLLGALVTAVI  
LVTGSVLVILENVTKILHPQPVNDEGILWLGIIAITINLLASLVVGKGTKNESILSLHFLEDTLGWVAVILMAIVLRFTDWYILDPL  
SLVISFFILSKALPRFWSTLKIFLDAVPEGLDIQVKSGLERLDNVASLNQLNLWTMDALEKNAIVHVCLKEMEHEMETCKESIRIFL  
KDCGFQNTITIEIDADLETHQTHKRKVCDLERSYEHQHZ

(SEQ. ID. NO. 83)

MIEYKNVALRYTEKDVLRDVLNQIEDGEFMYLVGPGSGKTTMLKMINRLEPTDGNIMYMDGKRIKDYDERELRLSTGYVLQAIA  
LFPNLTVAENIALIPEMKGWSKEEITKKTEELLAQVGLPVAEYGHRLPSELSGGEQQRVGIVRAMIGQPKIFLMDPEPSALDAISRK  
QLQVLTKELHKEFGMTTIFVTHDTEALKLADRIAQLDGEIRQVANPETILKAPATDFVADLFGGSVHDZ

(SEQ. ID. NO. 84)

MSAVAISAMTKVMQETHGNPSSIHGHRQAGKLLREARQELAQLLRTKPKHIFTSGGTEGNNTTIIGYCLRHQEQGKHIITTAIEH  
HAVLETIDYLVQHFGFEATHIQPENQEITAQQIQKALRDDTILVSTMFVNNETGNLLPIAEIGQILKQHPAAYHVDVQAIGKIPHSE  
ELGIDFLTASAHKHFHGPKGIGFLYASSMDFDSYLHGGDQEKKRAGTENLPAIVGMVAALKEDLEKQEEHFQHVQNLETAFLAEL  
EGIQYYLNRGKHHLPYVLNIGFPGQKNDLLLRDLAGISISGSACTAGVVQSSHVLEAMYGANSERLKESSLRISLPQNTVEDLQ  
TLAKTLKEIIGGZ

(SEQ. ID. NO. 85)

MLFKLSKEKIELGSLRSPARRIFLSFALVILLGSLLLSLPFVQVESSRATYFDHLFTAVSAVCVTGLSTLPVAHTYNIWGQIICLLLIQ  
IGGLGLMTFIGVFIYQSKQKLSLRSRATIQDSFSYGETRSLRKVFYSIFLTTFLVESLGAILLSFRILPQLGWGRGLFSSIFLAISAFCA  
GFDNLGSTSLFAFQTDLLVNLVIAGLIITGGLGFMVWFDLAGHVGRKKKGRLHFHTKLVLVLLTIGLLLFGTATTLFLEWNNAGTIG  
NLPVADKVLVSFFQVTVMRTAGFSTIDYTQAHPTLLIYILQMFLGGAPGGTAGGLKITTFFVLLVFARSELLGLPHANVARRTIAP  
RTVQKSFVFIIFLMSFLIGLLGITAKGNPPFIHLVFETISALSTVGVTANLTPDLGKLALSVIMPLMFMGRIGPLTLFVSLADYHPE  
KKDMIHYMKADISIGZ

(SEQ. ID. NO. 86)

MSDRITIGILGLGIFGSSVLAALAKQDMNIIAIDHAERINQFEPVLARGVIGDITDEELLRSAGIDTCDTVVVATGENLESSVLAVMH  
CKSLGVPTVIAKVKSQTAKKVLEKIGADSVISPEYEMGQSLAQTLFHNSVDVFQLDKNVSIVEMKIPQSWAGQSLSKLDLRGKYN  
LNLGFRREQENSPLDVEFGPDDLLKADTYILAVINNQYLDTLVALNSZ

(SEQ. ID. NO. 87)

MKLLSIAISSYNAAYLHYCVESLVIGGEQVVGILIINDGSQDQTQIEAECLASKYPNIVRAIYQENKCHGGA VNRGLVEASGRYFKV  
VDSDDWVDPRAYLKILETLQELESKGQEVDFVTNFVYEKQSRKKSMSYDSVLPVRQIFGWDQVGNFSKGQYTMHSLIYRT  
DLLRASQFZ

(SEQ. ID. NO. 88)

MKFNPQRYTRWSIRRLSVGVASVVVASGFFVLVGQPSSVRADGLNPTPGQVLPEETSGTKEGDLSEKPGDVTLTQAKPEGVTGN  
TNSLPTPTERTEVSEETSPSSLDTLFEKDEEAQKNPELTDVLKETVDTADVDGTQASPAETTPQVKGKVKENTKDSIDVPAAYLEK  
AEGKGPFTAGVNVQVPIYELFAGDGMLTRLLLKASDNAPWSDNGTAKNPALPPEGLTKGKYFYEVDLNGNTVGKQGQALIDQLR  
ANGTQTYKATVKVYGNKDGKADLTNLVATKNVDININGLVAKETVQKAVADNVKDSIDVPAAYLEKAKGEGPFTAGVNVHPIY  
ELFAGDGMLTRLLLKASDKAPWSDNGDAKNPALSPLGENVKTGQYFYQVALDGNVAGKEKQALIDQFRANGTQYTSATVNVY  
GNKDGKPDLDNIVATKVTININGLISKETVQKAVADNVKDSIDVPAAYLEKAKGEGPFTAGVNVHPIYELFAGDGMLTRLLLK  
SDKAPWSDNGDAKNPALSPLGENVKTGQYFYQVALDGNVAGKEKQALIDQFRANGTQYTSATVNVYGNKDGKPDLDNIVATK  
KVTININGLISKETVQKAVADNVKTVSMFQQPTZ

(SEQ. ID. NO. 89)

MKLKSYILVGYIISTLLTILVVFVAVQKMLIAKGEIYFLLGMTIVASLVGAGISLFLLLPVFTSLGKLKEHAKRVAAKDFPSNLEVQG  
PVEFQQLGQTFNEMSHDLQVSFDSLEESEREKGLMIAQLSHDIKTPITSIQATVEGILDGIIKESEQAHYLATIGRQTERLNKLVEELN  
FLTNTARNQVETTSKDSIFLDKLLIECMSEFQFLIEQERRDVHLQVIPESARIEGDYAKLSRILVNLVDNAFKYSAPGKLEVVAKL  
EKDQLSISVTDEQGQIAPEDLENIFKRLYRVETSRNMKTGGHGLGLAIAARELAHQLGGEITVSSQYGLGSTFTLVNLSGSENKAZ

(SEQ. ID. NO. 90)

MFGQTAQHGLTNSLKDWFIFLLNIGPQLAFFCQMLRCSRVEQGTGNHRRREFNMIQQIFSHFGMTHLGQIKLVYQESIDLELLVNA  
LNHHLLIDRLVLTNPQITIEIDRQIVHGLDLLKGRKDEKIIDIKSMFRQLELASTQQICPINQRVHHGILAFGEISDLVPAKNLPNRQDZ

(SEQ. ID. NO. 91)

MEHLATYFSTYGGAFFAALGIVLAVGLSGMGSAYGVGKAGQSAAALLKEQPEKFASALILQLLPGTQGLYGFVIGILIWLQLTPEL  
PLEKGVAYFFVALPIAIVGYFSAKHQGNVAVAGMQILAKRPKEFMKGAILAAMVETYAILAFVVSFILTLRVZ

(SEQ. ID. NO. 92)

MLKSEKQSRYQMLNEELSFLLEGETNVLANLSNASALIKSRFPNTVFAGFYLFDGKELVLGPFQGGVSCIRIALGKGVCGEAAHFQ  
ETVIVGDVTTYLNYISCDLAKSEIVVPMKNGQLLGVLDLDSSEIEDYDAMDRDYLEQFVAILLEKTAWDFTMFEEKSZ

(SEQ. ID. NO. 93)

MSVLEIKDLHVEIEGKEILKGVNLTCLKTEIAAIMGPNGTGKSTLSAAIMGNPNYEVTKEVLFDFGVNILELEVDERARMGLFLAM  
QYPSEIPGITNAEFLRAAMNAGKEDDEKISVREFITLDEKMELLMKEEMAERYLNEGFSGGEKKRNEILQLLMLEPTFALLDEID  
SGLDIDALKVSVKGVNAMRGEFGAMIITHYQRLNLYITPDVVHVMMEGRVVLSSGPELAARLEREGYAKLAEELGYDYKEELZ

(SEQ. ID. NO. 94)

MPYKRQRSFMSALSKLDSLYMAVVADHSKNPHHQKLEDAEQISLNNPTCGDVINLSVKFDAEDRLEDIAFLNSGCTISTASASM  
MTDAVLGKTKQEILELATIFSEMVCQKQDERQDQLGDAAFLSGVAKFPQRIKCATLAWNALKKTIENQEKQZ

(SEQ. ID. NO. 95)

MKIQDLRLKDVMLLDLQATEKTAVIDEMIKNLTDHGYVTDFTFKEGILAREALTSTGLGDGIAMPHSKNAAVKEATVLFASKNK  
GVDYESLDGQATDLFFMIAAPEGANDTHLAALAELSQYLMKDGFAKLRQATSADQVIELFDQASEKTEELVQAPANDSGDFIVA  
VTACTTGIAHTYMAQEALQKVAEMGVGIKVTETNGASGVGNQLTAEDIRKAKAIIIAADKAVEMDRFDGKPLINRPVADGIRKTE  
ELINLALSGDTEVYRAANGAKAATASNEKQSLGGALYKHLMSGVSQMLPFVIGGGIMIALAFLIDGALGVNENLGNLGSYHELA  
SMFMKIGGAAGFLMLPVFAGYVAYSIAEKPLVAGFVAGAIAGEGAFGKIPYAAGGEATSTLAGVSSGFLGALVGGFIAGALVL  
AIKKYVKVPRSEGAKSILLPLLTGTILTFVMLAVNIPMAAINTAMNDFLGGGLGGGSAVLLGIVLGGMMAVDMGGPVNKAAYV  
FGTGLAATVSSGGSVAMAAVMAGGMVPLAIFVATLLFKDKFTKEERNSGLTNIIMGLSFITEGAIPFGAADPARAIPSFILGSAVA  
GGLVGLTGIKLMAHPGGIFVIALTSNALLYLVSVLVGAIVSGVVYGYLRKPQAZ

(SEQ. ID. NO. 96)

MANKNTSTTRRRPSKAELERKEAIQRMLISLGIAILLIFAFLGAAGITLYNLIRLLVGSLEYLAIFGLLIYFFFKWIRKQEGLLSGF  
FTIFAGLLIFEAYLVWVKYGLDKSVLKGTMQVVTDLTGFRFTTSFAGGGLIGVALYIPTAFLFSNIGTYFIGSILVGSLLVSPWSVY  
DIAEFFSRGFAKWEGHERRKEERFVKQEEKARQKAKEARLEQEETEKALLDLPPVDMETGEILTEEAVQNLPIPEEKWVEPEII  
LPQAEKPFQEEDSDDEDVQVDFSAKEALEYKLPQLFAPDKPKDQSKKEKKIVRENIKILEATFASFGIKVTVERAIGPSVTKEYE  
VKPAVGVRVNRISNLSDDLALALAAKDVRIEAPIPGKSLIGIEVPNSDIATVSFRELWEQSQTAKENFLEIPLGKAVNGTARAFDLSK  
MPHLLVAGSTGSGKSVAVNGIIASILMKARPQVKFMMVDPKMWELSVYNDIPHLLIPVVTNPRKASKALQKVVDENENRYELFA  
KVGVRNIAGFNAKVEEFNSQSEYKQIPLPFIVVIVDELADLMMVASKEVEDAIIRLGQKARAAGIHMILATQRPVSDVISGLIKANV  
PSRVAFVSSGTDSTRILTDENGAEKLLGRGDMLFKPIDENHPVRLQGSFISDDDDVERIVNFIKTQADADYDESFDPGEVSENEGEFS  
DGDAGGDPLFEEAKSLVIETQKASASMIQRRLSVGFNRATRLMEELEIAGVIGPAEGTKPRKVLQZZ

(SEQ. ID. NO. 97)

MSYFKKYKFDKSQFKLGMRTFKTGIAVFLVLLIFGFFGWKGLQIGALTAVFSLRESFDESVDHFGTSRILGNSIGGLYALVFFLLNTFF  
HEAFWVTLVVPICTMLTIMTNVAMNNKAGVIGGVAAMLITLSPSGETILYVVRVLETFMGVFVAIIVNYDIDIRLFLFLEKKEKZ

(SEQ. ID. NO. 98)

MNKSEHRHQLIRALITKNKIHTQAEQLALLAENDIQVTQATLSRDIKNMNLKSVREEDSAYVVLNNGSISKWEKRLELYMEDALV  
WMRPVQHQLVLLKTLPLGLAQSFSGIIDLTSFPDAIATLCGNDVCLICEDADTAQKCFEELKKFAPFFFFEEZ

(SEQ. ID. NO. 99)

MKSIKLNALSYMGIRVLNIIFFILTGTYYARVLDRTDYGYFNSVDLTLSFFLPFATYGVYNYGLRAISNVKDNKKDLNRTFSSLFYLC  
IACTILTAVYILAYPLFFTDNPIVKVYLVMGIIQIAQIFSIEWVNEALENYSFLFYKTAIRILMLVSIFLKVNEHDIVVYTLVMSL  
STLINYLISYFWIKRDIKLVKIHLSDFKPLFLPTAMLVFANANMLFTFLDRFLVKTGIDVNVSYTTIAQRIVTVIAGVVTGAIGVS  
PRLSYLGGKGDKEAYVSLVNRGSRIFNFFIPLSFGMLVLPNAILLYGSEKYIGGGILTSLFAFRTHIALDITLGSQILFTNGYEKRIT  
VYTVFAGLLNLGLNSLLFFNHIVAPEYLLTMTLSETSLVFIIFHRRKQLIHLGHIFSITYVRYSLSLSFVAIYFLINFPVDMVIN  
LPFLINTGLIVLLSAISYISLLVFTKDSIFYEFLNHVLALKNKFKKKSZ

(SEQ. ID. NO. 100)

MELFMKITNYEIKLKKSGLTNQILKVLEYGENVDQELLGDIADISGCRNPVFMERYFQIDDAHLSKEFQKFPFSILDCCYPW  
DLSEIYDAPVLLFYKGNLDLLKFPKVAVVGSRACSKQGAKSVEKVIQGLLENELVIVSGLAKGIDTAHMAALQNGGKTAIVIGTGL  
DVFPKANKRLQDYIGNDHLVLSEYGPGEQPLKFHFPARNRIIAGLCRGVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLS  
DGCHHLIQEGAKLVTSGQDVLAEEFZ

(SEQ. ID. NO. 101)

MKQLTVEDAKQIELEILDYIDTLCKKHNNIYIINYGTLLIGAVRHEGFIPWDDDDIDLSMPREDYQRFINIFQKEKSKYKLLSLETDKNY  
FNNFIKITDSTTKIIDTRNTKTYESGIFIDIFIDRFDPPKVIDTCYKLESFKLLSFSKHKNIVYKDSLLKDWIRTAFWLLRPVSPRYFA  
NKIEKIQKYSRENGQYMAFIPSKFKEKEVFPSGTFDKTIDLPFENLSLPAPEKFDITLTQFYGDYMTLPPEEKRFYSHEFHAYKLED  
Z

(SEQ. ID. NO. 102)

MIKINHLTITQNKDLRLVSDLTMTIQDGEKVAIIEEGNGKSTLLKILMGEALSDFTIKGNISQSDYQSLAYIPQKVPEDLKKKTLHD  
YFFLDSDLDYSILYRLAEELHFDNSRNASDQEIENLSGGEALKIQLIHELAKPFEILFLDEPSNDLDLETVDWLKGGQIKTRQTVIFIS  
HDEDFLSETADTIVHLRLVKHKEAETLVEHLDYDSYSEQRKANFAKQSQQAANNQRAYDKTMEKHRRVKQNVETALRATKDS  
TAGRLLAKMKTKVLSQEKRYEKAQSMQKPLEEEQIQLFFSDIQLPASKVLVQLEKENLSIDDRVLVQKLQTLTVRGQEKIGIIGP  
NGVGKSTLLAKLQRLNLDKREISLGFMQDYHKKLQDLSPAIYLSKTGEKEELQKIQSHLASLNFSYPQMQRSLSGGQGGKL  
LLDLVLKPNFLLDDEPTRNFSPTSQPQIRKLFATYPGGLITVSHDRRFLKEVCSIIYRMTEHGLKLVNLEDLZ

(SEQ. ID. NO. 103)

MKPKTFYNLLAEQNLPLSDQKQEQERYFELLVEWNEKINLTAITDKEEVYLLKHFYDSIAPILQGLIPNETIKLLDIGAGGFPSPM  
KILYPELDVTIISLNLKRINFLQLLAQELDLNGVHFYHGRAEDFAQDKNFRAQYDFVTARAVARMQVLSLTIPYLVGGKLLALK  
ASNAPELLEAKNALNLLFSKVEDNLSYALPNRDPRIYTVVEKKKETPNKYPRKAGMPNKRPLZ

(SEQ. ID. NO. 104)

MSIKLIAVDIDGTLVNSQKEITPEVFSAIQDAKEAGVKVVIATGRPIAGVAKLLDDLQLRDEGDYVVTFNAGLVQETATGHEIISL  
TYEDYLDMEFLSRKLGVMHMAITKDGITYANRNIGKYTVHESTLVSMPIFYRTPEEMAGKEIVKCMFIDEPEILDAAIEKIPAEFYER

YSINKSAPFYLELLKKNVDKGSATHLAELKGLTKDETMAIGDEENDRAMLEVVGPNVVMENGNEPKIKIAYITKTNDSESGVAH  
AIRTWVLZ

(SEQ. ID. NO. 105)

MTWILGVIALIVFVIVSYNGLVKNRMQTKEAWSQIDVQLKRRNDLLPNLIETVKGYAKYEGSTLEKVAELRNQVAAATSPAEM  
KASDALTRQVSGIFAVAESYDLKASANFVKLQEELTNTENKISYSRQLYNSVSNYNVKLETFFPSNIIAGMFGFKAADFLQTPEEE  
KSPVKVDFSGLDZ

(SEQ. ID. NO. 106)

MLFDQIASNKRKTWILLVFFLLALVGYAVGYLFIRSGLGGLVIALIIGFIYALSMIFQSTEIVMSMNGAREVDEQTAPDLYHVVE  
DMALVAQIPMPRVFIIDDPALNAFATGSNPQNAAVAATSGLLAIMNREELEAVMGHEVSHIRNYDIRISTIAVALASAITMLSSMAG  
RMMWWGGAGRRRSDDDRDGNLEIIMLVSLLAIVLAPLAATLVQLAISRQREFLADASSVELTRNPQGMINALDKLDNSKPM  
RHVDDASSALYINDPKKGGGFQKLFYTHPPISERIERLKQMZ

(SEQ. ID. NO. 107)

MKLNIEIRKQSEGLNFEQTLDDLVDLARNQEILDVKDILAVGKVQYEDRMFLDYQLSYTIVLASSRSMPELVESYPVTEVF  
MEGATNQLDQEVLDLDDLVLPENGELDLAESVSDNILLNIPIKVLTAEEEEAGQGFISGNDWQIMTEEEYQAQKAVKKEENSFPAGL  
QGLFDGDEZ

(SEQ. ID. NO. 108)

MKRQALVVFSGGQDSTTCLFWVMQHYETVEAVTFAYGQRHHEIQITREIAKEQGIRHHILDMSLGQITAQPDFATIHISYIPDK  
LCVESKSLKLYFSYRNHGDFFHENCINTIGKDLVNLLDPRYLEVWGKFTPRGGISIDPYNYGKQGTKYEGLAEQRLFQHDLYPEK  
IDNRZ

(SEQ. ID. NO. 109)

MTETVEDKVSHSITGLDILKGIVAAGAVISGTVATQTKVFTNESAVLEKTVEKTDALATNDTVVLGTISTSNSASSTSLSASESASTS  
ASESASTSASTSASESASTSASTSISASSTVVGSTAAATEATAKKVEEDRKKPASDYVASVTNVNLQSYAKRRKRSVDSIEQ  
LLASIKNAAVFSGNTIVNGAPAINASLNAIKSETKVYTGEVDSVYRVPYIYKLVKTNDGSKLTFTYTVTYVNPKTNDLGNISSMRP  
GYSIYNSGTSTQMTLTGSDLGKPSGVKNYITDKNGRQVLSYNTSTMTTQSGGYTWGNGAQMNNGFAKKGYGLTSSWTVPITGT  
DTSFTFTPYAARTDRIGINYFNGGGKVVESSTTSQSLSQSKLSVSASQSASASASTSASASASTSASASASTSASASASTSASVSAST  
SASASASTSASASASTSASESASTSASASASTSASASASTSASASASTSASESASTSASASASTSASESASTSASASASTSASASASTS  
SGSASTSTSASASTSASASASTSASASASISASESASTSASESASTSTSASASTSASESASTSASASASTSASASASTSASASASTSASAS  
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QRQPVRQLQQVPVLQSQHQVLPQPHRQVPRQLQQAQHLNQRQAPQLQQVPVRQPQRQVRQPQQVVLVHQLQHQRVHRLRR  
QPVHQSQQVPVRQLPHQQVPRQLQAPVRRQLQQVLAPQPQPVRQPQQVSRQLNRHQVRPLQQVLAPQPQRQQVHRLQRQRV  
RLNRHQVRPLQQVLAPQPQRQQVHRLQHQRVRPLQQVLAPQPQRQQVHRLQRQRVRLSQHQVRQPQAQHLNQLHQPVRQP  
QHRQAPQLQQVPVRQPQRQVRRLQQVPVRQPQQVPVRQPQRQVRPQVHLNRHQPVRRPQQVVLVHQLQHQRVHRLQHQPVR  
HQSQQVPVRQFRINKCLGFSKYZ

(SEQ. ID. NO. 110)

MGVETWFYSSICWLAIGLSVWKFPMYMTAANGGGGFLIFLISTILIGFPLLLAEFALGRSAGVSAIKTFGKLGKNNKYNFIGWIGA  
FALFILLSFYFVIGGWILVYLGIEFGKLFQLGGTG DY AQLFTSIIISNPAIALGAQAIFILLNIFIVSRGVQKGIERASKVMMPLLFIVFV  
FIGRSLSLPNAMEGVLYFLKPDFSKLTSTGLLYALGQSFFALSLGVTVMILTYASYLDKKTNLVQSGISIVAMNISISIMAGLAIFQAR  
SPFNIQSEGGPSLLFIVLPQLFDKMPFGTIFYVLFLLFLFATVTFSSVMLEINVDNITNQDNSKRAKWSVILGILTFVFGIPSALS YGV  
MADVHIFGKTFDFAMDVLVSNLLMPFGALYLSLFTGYIFKKALAMEELHLDERAWKQGLFQVWFLFLRFFVSSSQSSSLWSSLPNL  
CNQKGLEZ

(SEQ. ID. NO. 111)

MLKKWQLKDVILLAFLSIFFGGVFGVSGYVYNILSLLTPLGLQAFANEILFGLWCMAPIAAIFVPRVGSATIGEVLAAAEVLYG  
SQFGLGALLSGFVQGLGSEFGFIVTKNRYESWLSLTANSIGITLVSFVYIEYIKLGYAFSLPFVLSLLVVRFSISVYFFCTILVRAIVKL  
YHQFATGGKAZ

(SEQ. ID. NO. 112)

MVKVATQTPHISLFLILSLETSPISIALTSLVVAFCILFMYLRYRRFKMLAWMIILAILPSFANYWAVQLHGDASQAVMLGTRAFVT  
VCIGLVFVSSVSLKELLLYLAQKGLSRWSYALIVVFNFPFIQEQEIKSLKEACLLRGQELHFWSPLIYSKVLMTVFRWRHLYLRLAL  
SAHG YDEHAQLKNSYRTFYIPKTKLIYLLFFLLQTSLSFLZ

(SEQ. ID. NO. 113)

MRKHQLQVHKLTILSMMIALDVVLTPIFRIEGMAPMSSVVNLAGIMMGVYALAMATVTA FIRM T T Q G I P P L A T G A T F G A L L A G  
LFYKYGRKFHYSALGEILGTGIISIVSYPMVLF TGSAAKLSWFIYTPRFFGATLIGTAISFIAFRFLIKQEFFFKKVQGYFFSERIDZ

(SEQ. ID. NO. 114)

MQEFTNPFPIGSSSLIHCITNEISCEMLANGILALGCKPVMADDSREVLDFTKQSQUALFINLGHLSAEKEKAIRMAASYANQSSSLPMV  
VDAGVITSSIRKSLVKDLDYRPTVLKGNMSEIRSLVGLKHHGVGDASAKDQETEDLLQVLKDWQCQTYPGMSFLVTGPKDLV  
VSKNQVAVLGNCTELDWITGTGDLVGALTAVFLSQGKTGFASCLAVSYLNIAAEKIVVQGMGLEEFYQVVLNQLSLLRRDEN  
WLDTIKGEVYEZ

(SEQ. ID. NO. 115)

MNHKIAILSDVHGNATALEAVIADAKNQGASEYWLLGDIFLPGPGANDLVALLKDLPTASVRGNWDDRVLALDGGYGLEDPQ  
EVQLLRMTQYLMERMDPATIVWLRSLPLEKKEIDGLRFSISHNLPDKNYGGDLLVENDTEKFDQLLDAETDVAVYGHVHKQLL  
RYGSQQQIINPGSIGMPYFNWEALKNHRSYAVIEVEDGELLNIQFRKVA YDYEAELAKSKGLPFIEMYEELRRDDNYQGHNL  
ELLASLIEKHGYVEDVKNFFDFLZ

(SEQ. ID. NO. 116)

MNVNQIVRIIPTLKANNRKLNETFYIETLGMKALLEESAFLSLGDOTGLEKLVEEAPSMRTRKVEGRKKLARLIVKVENPLEIEGIL  
SKTDSIHRLYKQGNGYAFEIFSPEDDLILHAEDDIASLVEVGEKPEFQTDLASISLSKFEISMELHLPTDIESFLESSEIGASLDFIPAQG  
QDLTVDNVTWDLMLKFLVNELDIASLRQKFESTEIFPKSEKFFLGKDRNNVELWFEVZ

(SEQ. ID. NO. 117)

MKWTIKIIEEIEAGIYPGASFAYFKDNQWTEFYLGQSDPEHGLQTEAGLVYDLASVSKVVGVTCTFLWEIGQLDIDRLVID  
FLPESDYPDITIRQLLTHATDLDPFIPNRDLLTAPELKEAMFHLNRRSQPAFLYSDVHFLLLGFLERIFNQDLVDILKDQVWKPWGM  
TETKFGPVELAVPTVRGVEAGIVHDPKARLLGRHAGSAGLFSTIKDLQIFLEHYLADDFARDLNNQNSPLDDKERSLAWNLEGDW  
LDHTGYTGTFIMWNRQKQEAIFLSNRTYEKDERAQWILDRNQVMNLIRKEEZ

(SEQ. ID. NO. 118)

MMKKTYNHILVWGVIFYSICIVCFCTPQEOSTVGVTGPIQHLGRLVFLLPFNSLWKLGEVSDIGQLCWIFLQNILNVFLFFLIFQ  
LLYLPNLRKTKKVLFSFLVSLGIECTQLILDFFFDFNRVFEIDDLWTNTLGGYLAWLLYKRLHKNKVRNZ

(SEQ. ID. NO. 119)

MKPIILLTFARHKFYVLLTLLFLALVYRDVLMTYFFFDIHAPDLAKFDGQAIKNDLLKSALDFRILQFNLGFYQSFIIPIIIVLLGFQYI  
ELKNKVLRLSIGREVSQGLKRLTLQVASIPCLIIYLVTLIIAITYFFGTFSPLGWSLSDGSGLRLLDGEIKSYLFFTCVLLIGIF  
INAIYFLQIVDYVGNVTRSAITYLMLFLWLGSMILYSALPYMVPMTSLMQASYGVSMLKLFPTYIYIVPYMVLEKYEDNVZ

(SEQ. ID. NO. 120)

MFKVLQKVKGAFMLPIAILPAAGLLLGIGGALSNTPTTIATYPILDNSIFQSIFQVMSSAGEVVSFNSLULLCVGLCIGLAKRDKGTAA  
LAGVTGYLVMTATIKALVKLFMAEGSAIDTGVIGALVVGIVAVYLHNRYNNIQLPSALGFFGSGSRFVPIVTSFSSILIGVFFVWPPF  
QQLLVSTGGYISQAGPIGTFLYGLMRLSGAVGLHHIYPMFWYTELGGVETVAGQTVVGAQKIFFAQLADLAHSGLFTGTRFFA  
GRFSTMMFGLPAACLAMYSVPKNRRKKYAGLFFGVALTSTFITEPIEFMFLFVSPVLYVHAFLDGVSFADVLNISIGNTFSG  
GVIDFTFLGILQGNAKTNWVLQIPFGLIWSVLYYIIFRWFTTQFNVLTTPGRGEEVDSKEISESADSTSNTADYLYKQDSLQIIRALGGSN  
NIEDVDACVTRLRVAVKEVNQVDKALLKQIGAVDVLEVKGQIAYGAKAILYKNSINEILGVDDZ

(SEQ. ID. NO. 121)

MKFRKLACTVLAGAAVLGLAACGNSGGSKDAAKSGGDGAKEITWWAFPVFTQEKTDGSGVGTYEKSIIEAFEKANPDIKVKLETI  
DFKSGPEKITTAIEAGTAPDVLFDAPGRIIYQYKNGKLAELNDLFTDEFVKDVNNENIVQASKAGDKAYMYPISAPFYMAMNKK  
MLEDAGVANLVKEGWTTDDFEKVLKALKDKGYTPGSLFSSGQGGDQGTAFISNLYSGSVTDEKVSKEYTTDDPKFVKGLEKATS  
WIKDNLINNGSQFDGGADIQNFANGQTSYITLWAPAQNGIQAKLLEASKVEVVEVPFSPDEGKPALEYLVNGFAVFNNKDDKKVA  
ASKKFIQFIADDKWGPDKDVRTGAFVPTSFGKLYEDKRMETISGWTQYYSPTYNTIDGFAEMRTLWFPMLQSVSNGDEKPADA  
LKAFTKANETIKKAMQZ

(SEQ. ID. NO. 122)

MQSTKKPLTAFTVISTIILLTTLVLFIFPFYWILTGAFKSQPDITVIPPQWFPKMPTMENFQQLMVQNPALQWMWNSVFISLVTMFL  
VCATSSLAGYVLAKKRFYQORILFAIFIAAMALPKQVVLVPLVRIVNFMGIHDTLWAVILPLIGWPFVGLMKQFSENIPTELLESA  
KIDGCGEIRTFWSVAFPIVKPGFAALAIPTFINTWNYFMQLVMLTSRNNLTISLGVATMQAEMATNYGLIMAGAALAAVPIVTVF  
LVFQKSFTQGITMGAVKGZ

(SEQ. ID. NO. 123)

MKIMFKNFNILLNRKIVLLLRIVLMMILINHLLSTAVQKQDAVIFFKRELISIFSNDYSEANLEIPKLLLNLSLFMVGWLSVILLES  
DLADHYHLLIRYQSSSFDDYTRKRLVVISKFQDLFWFLGLLPLGIHFKTVALFFLLAQLMMLYLLLSYLIALLISAGAGFSFFLYF  
LAFVQGQEWMDHIVTVYLVLLSLLVMLIVSRLEKFKKGZ

(SEQ. ID. NO. 124)

MGKGEMGKGVIGLEFDSVLVNKAPTQLANGKTATFLTQYDSKTLTFAVDKEDIGQEIIAGKGSIESMHNLPVNLGARVPGGV  
NGSKAAVHEVPEFTGGVNGTEPAVHEIAEYKGSDSLVTLTTKKDYTYKAPLAQQALPETGNKESDLLASLGLTAFFLGLFTLGKK  
REQZ

(SEQ. ID. NO. 125)

MKKTFFLLVLGLFCLLPLSVFAIDFKINSYQGDLYIHADNTAEFRQKIVYQFEEDFKGQIVGLGRAGKMPSGFDIDPHPKIAAKNG  
AELADVTSEVTEEADGYTVRYNPGQEGDIVEVDLVWNLKNLLFLYDDIAELNWQPLTDSSESIEKFEFHVVRGDKGAELFFHTG  
KLFREGTIEKSNLDYTIRLDNLPKRGVELHAYWPRTDASARDQGLKGNRLEEFNKIEDSIVREKDQSKQLVTWVLPISILSLLLS  
VCFYFIYRRKTTSPVKYAKNHRLYEPPMELEPMVLSEAVYSTSLEEVSPLVKGAGKFTFDQLIQATLLDVIDRGNVSIIEGDAVGL  
RLVKEDGLSSFEKDCNLAFSGKKEETLSNLFADYKVSDSLRYRAKVSDEKRIQARGLQLKSSFEVVLNQMQEGVVRKRVSWGLP  
DYRPLTGGEKALQVGMGALTILPLFIGFLFLYSLDVHGYLYLPLPILGFLGLVLSVFYWKLRDNRDGVLEAGAEVYYLWT  
SFENMLREIARLDQAELESIVVWNRLLVYATLFGYADKVSMLMKVHQIQVENPDINLYVAYGWHSTFYHSTAQMASHYASVANTA  
STYSVSSGSGSSGGGFGSGGGGSGSIFAFZ

(SEQ. ID. NO. 126)

MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVFELKNNTDGTTSQRTEAQTGEAIFSNIK  
PGTYTLTEAQPPVGYKPSSTKQWTVVEKNGRTTVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNY  
ERVIPEGTLISKRIYQVNNLDDNQYGIELTVSGKTVYEQDKSVPLDVILLDNSMSNSNIRNKNARRAERAGEATRSIDKITSDE  
NRVALVYASTIFDGTFTVEKGVADKNGKRLNDSLFWNYDQTSFTNTKDYSYLKLTNDKNDIVELKNKVPTAEHDHGNRLM  
YQFGATFTQKALMKADEILTQARQNSQKVIFHITDGVPMTSPINFNHATFAPSQYNQLNAFFSKSPNKDGILLSDFITQATSGEH  
TIVRGDQSYQMFTDKTVYEKGAPAAFPVKPEKYSEMKAAGYAVIGDPINGGYIWLNWRESILAYPFNSNTAKITNHGDPTRWYY  
NGNIAPDGYDVFTVVGINGDPGTDEATATSMQSISSKPENYTNVTDTTKILEQLNRYFHTIVTEKKSIENTITDPMGELIDLQGT  
DGRFDPADYTLTANDGSRLENGQAVGGPQNDGGLKNKVLVYDTTEKIRVTGLYLGTEKVTLTYNVRLNDEFVSNKFYDTNG  
RTTLHPKEVEQNTVRDFPIPKIRDPVKYPEITISKEKLGDIIEFIKVNKNDDKKPLRGAVFSLQKQHPDYPDIYGAIQNGTYQNVRT  
GEDGKLTFFKNLSDGKYRLFENSEPAGYKPVQNKPIVAFQIVNGEVRDVTISVPQDIPAGYEFTNDKHYITNEPIPPKREYPRTGIG  
MLPFYLIGCMGMMGGVLLYTRKHPZ

(SEQ. ID. NO. 127)

MKSINKFLTMLAALLLTASSLFSAA TVFAAGTTTTSVTVHKLLATDGDMDKIANELETGNYAGNKVGVLPANAKEIAGVMFVWT  
NTNNEIHDENGQTLGVNIDPQTFKLSGAMPATAMKKLTEAEGAKFNTANLPAKYKIYIEHSLSTYVGEDGATLTGSKAVPIEIELP  
LNDVVDAAHVYPKNTAKPKIDKDFGKANPDTPRVDDKTPVNHQVGDVVEYIEIVTKIPALANYATANWSDRMTEGLAFNKGTV  
KVTVDVDALEAGDYALTEVATGFDLKLTDAGLAKVNDQNAEKTVKITYSATLNDKAIVEVPESNDVTFNYGNNPDHGNTPKPNK  
PNENGDLTLTKTWV DATGAPIAGAEATFDLVNAQTGKV VQTVTLTTDKNTVTVNGLDKNTYKFFVERSIKGY SADYQEITTAG  
IAVKNWKDENPKPLDPTEPKVV TYGKKFVKVNDKDNRLAGAEFVIANADNAGQYLARKADKVSQEEKQLVVTTKDALDRAVA  
AYNALTAQQQTQKEKVDKAQAAYNAAVIAANNAFEWVADKDNENVVKLVSDAQGRFEITGLLAGTYYLEETKQPAGYALL  
TSRQKFVETATSYSATGQGIEYTAGSGKDDATKVVNKKITPQTGGIGITIIFAVAGAAIMGIAVYAYVKNKDEDQLAZ

(SEQ. ID. NO. 128)

MTMQKMQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEVVSQLPSRDGHRQLQVWKLDDSSYSYDDRQVIVRDLHS  
WDENKLSSFKKTSEFMTFLENQIEVSHIPNGLYYVRSIIQTDVSYPAEFLFEMTDQTV EPLVIVAKKTD TMTTKVKKLKVDDQDHNH  
LEGVGFKLVS VARDVSEKEVPLIGEYRYS SSGQVGR TLYTDKNGEIFVTNLPLGNYR FKEVEPLAGYAVTTLD TDVQLVDHQLVTI  
TVVNQKLPRGNVDFMKVDGRNTSLQGAMFKVMKEESGHYTPVLQNGKEVVTS GKDGRFRVEGLE YGTYLWELQAPTGYV  
QLTSPVSFTIGKDRKELVTVVKNKRPRIDVPDTGEETLVYLDACCHFVWVZ

(SEQ. ID. NO. 129)

MSHIYLSIFTSLLMLGLVNVAQADEYL RIGMEAA YAPFNWTDQDDSN GAVKIDGTNQYANGYDVQIAKKIAKDLGKEPLVVK  
KWEGLVPALTS GKIDMIIAGMSPTAERKQEI AFSSSYTSEPVLLVKKDSAYASAKSLDDFN GAKITSQQGVYLYNLIAQIPGAKKE  
TAMGDF AQMRQALEAGVIDAYV SERPEALTAEAN SKFKMIQVEPGFKTGEEDTAIAIGLRKNDNRISQINAS IETISKDDQV ALM  
DRMIKEQPAEATTTETSSSFFSQVAKILSENWQQLLRGAGITLLISIVGTIIGLIIGLAIGVFRTAPLSENKV IYGLQKLVGWVLNVYI  
EIFRGTPMIVQSMV IYGTAAQAFGINLDRTLAAIFIVSINTGAYMTEIVRGILAVDKGQFEAATALGMTHNQTMRKIVLPQVVRNI  
LPATGNEFVINIKDTSVLNVISVVELYFSGNTVATQTYQYFQTFTIIA VIYFVL TFTVTRILRFIERRMDMDTYTTGANQM QTEDLKZ

(SEQ. ID. NO. 130)

MTQAILEIKHLKKS YGQNEVLKDISLTVHKGEVISIIGSSSGSKSTFLRSINLLETPTDGGILYHGQNVLEKGYDLTQYREKLG MVFQ  
SFNLFENLNVLENTIVAQTTVLKRERTEAEKIAKENLEKVGMGERYWQAKPKQLSGGQKQRVAIARALSMNPDAILFDEPTSA LD  
PEMVGEVLKIMQDLAQEGLTMIVVTHEMEFARDVSHRVIFMDKG VIAEEGKPEDLFTNP KEDRTKEFLQRYLKZ

(SEQ. ID. NO. 131)

MKKYQLLFKISAVFSYLFFVFSLSQLTLIVQNYWQFSSQIGNLFWIQNILSLLFIGVMIVVLVKTGHGYLFRIPRKKWLWYSILT VLV  
LVFQISFNVTAKH VQSTAEGWAVLIGYSGTNFAELGIYIALFFLVPLMEELIYRGLLQHAF FKHRSRFGDLLLPSILFALPHFSSLP S  
LLDIFVFATVGIIFAGLTRYTKSIYPSYAVHVINNIVATFPFLLTFLHRVLGZ

(SEQ. ID. NO. 132)

MNKKQWLGLGLVAVAAVGLAACGNRSSRNAASSSDVKTKAAIVTDTGGVDDKSFNQSAWEGLQAWGKEHNLSKDNNGFTYFQS  
TSEADYANNLQQAAGSYNLIFGVGFALNNAVKDAAKEHTDLN YVLIDDVIKQKNVASVTFADNESGYLAGVAAAKTTKTKQV  
GFVGGIESEVISRFEAGFKAGVASVDPSIKVQVDYAGSFGDAAGKGTIAAAQYAAGADIVYQVAGGTGAGVFAEAKSLNESRPEN  
EKVWVIGVDRDQEAEGKYTSKD GKE SNFVLVSTLKQVGTTVKDISNKAERGEFP GGQVIVYSLKDKGVDLAVTNLSEEGKKA VE  
DAKAKILDG SVKVPEKZ

(SEQ. ID. NO. 133)

MSKKLQQISVPLISVFLGILLGAIVMWIFGYDAIWGYEELFYTA FGSRLRGIEIFRAMGPLVLIGLGF AVASRAGFFNVGLPGQALAG  
WILSGWFALSHPDMPRPLMILATIVIALIAGGIVGAIPGILRAYLGTSEVIVTIMMNYIVLYVGNAFIHA FPKDFMQSTDSTIRVGAN  
ATYQTPWLAELTGNSRMNIGIFFAIIA VAVIWFMLKKTTLGFEIRAVGLNPHASEYAGISAKRTIILSMIISGALAGLGGA VEGLGTF  
QNYYVQGSLSAIGFNGMAVSLLAANSPIGILFAAFLFGVLQVGAGPMNAAQVPSELVSIVTASIIFFVSVHYLIERFVKPKKQVKGG  
KZ

(SEQ. ID. NO. 134)

MGVKKKKLKLTSLLGLSLLIMTACATNGVTS DITAESADFW SKLVYFFAEIIRFLSFDISIGVG IILFTVLIRTVLLPVFQVQMVASRKM  
QEAQPRIKALREQYPGRDMESRTKLEQEMRKVFKEMGVRQSDSLWPILIQMPVILALFQALS RVDFLKTGHFLWINLGSDTTLVL  
PILAAVFTFLSTWLSNKA LSERNGATTAMMYGIPVLIFIAVYAPGGVALYWTVSNA YQVLQTYFLNNPFKIIAEREAVVQAQKDL  
ENRRKRKAKKKAQKTKZ

(SEQ. ID. NO. 135)

MVIDPFAINELDYLVSHFSDHIDPYTAAAILNNPKLEHV KFIGPYHCGRIWEGWGVPKERII VVKPGD TIELKDMKIHAVESFDR  
TCLVLPVNGADETG GELAGLAVTDEEMAQKAVNYIFETPGGTIYHGADSHFSNYFAKHGKDFKIDV ALNNYGENPVGIQDKMT  
SIDLLRMAENLR TKVIIPVHYDIWSNFMAS TNEILELWKMRKDR LQYDFHPFIWEVGGKYTPQDQHLVEYHHPRGFDDCFEQDS  
NIQFKALLZ

(SEQ. ID. NO. 136)

MFLSGWLSSFANTYIHDLLGVLPD SPFLNAFESAIAAPLVEEPLKLLSLVFLALIPVRKLKSLFLLGIASGLGFQMIKDIGYIRTDLP  
EGFDFTISRILERIISGIASHWTFSGLA VVG VYLLYRAYKGQKVGKKQGLIFLGLALGTHFLFNSPFVELETELPAIPVVTALYGF  
YHAYCFVEKHNE LMTZ

(SEQ. ID. NO. 137)

MKVEPRCDVLSRMSHFFIRILIMELQELVERSWAIRQAYHELEV KHHDSKWTVEEDLLALSNDIGNFQRLVMTKQGRYYDETPT Y  
LEQKLSENIWWLELSQRLDIDILTEMENFLSDKEKQLNVRTWKZ

(SEQ. ID. NO. 138)

MLDWKQFFLAYLRSRSLFIYLLSLAFLVLLFQFLFASLGIYFLYFFFLCCFVTILFFTWDILVETQVYRQELLYGEREAKSPLEIALA  
EKLEAREMELYQQRSAERKLTDLDDYYTLVWHQIKTPIAASQLLVAEVVDRQLKQQLQEIQFIDSYTNLVQLYLRLESFHDDL V

LKQVQIEDLVKEIRKYALFFIQKGLNVNLHDLDEIVTDKKWLLVVIEQIISNSLKYTKEGGLEIYMDDQELCIKDTGIGIKNSDVL  
RVFERGFSGYNGRLTQQSSGLGLYLSKKISEELGHQIRIESEVGKGTTVRIQFAQVNLVLEZ

(SEQ. ID. NO. 139)

MELNTHNAEILLSAANKSHYPQDELPEIALAGRSNVGKSSFINTMLNRKNLARTSGKPGKTQLLNFFNIDDKMRFVDVPGYGYAR  
VSKKEREKWGCMIEEYLTRENLRVSVSLVDRHDPSADDVQMYEFLKYEIPVIVATKADKIPRGKWNKHESAIKKKNLNDPS  
DDFILSSVSKAGMDEAWDAILEKLZ

(SEQ. ID. NO. 140)

MTKKQLHLVIVTGMGSGAGKTVAIQSFEDLGYFTIDNMPALLPKFLQLVEIKEDNPKLALVDMRSRSFFSEIQAVLDELENQDGL  
DFKILFLDAADKELVARYKETRRSHPLAADGRILDGIKLERELLAPLKNMSQNVVDTELTPRELKTLAEQFSDEQAQSFRIEV  
MSFGFKYGIPIDADLVFDRFLPNPYLPELRNQTGVDEPVYDVMNHPESEDFYQHLLALIEPILPSYQKEGKSVLTIAMGCTGGQ  
HRSVAFAKRLAQDLSKNWSVNEGHRDKDRRKETVNR5Z

(SEQ. ID. NO. 141)

MRKPKITVIGGGTGSPIVLKSLREKDVIEAIVTVADDGSSGELRKNMQQLTPPGDLRNVLVAMSDMPKFYEKVFQYRFSEDAG  
AFAGHPLGNLIIAGLSEMGGSTYNAMQLLSKFFHTTGKIYPSSDHPPLTHAVFQDGTVEVAGESHIVDHRGIIDNVYVTNALNDDTPL  
ASRRVVQTILESDMIVLGPGLFTSILPNIVIKEIGRALLETKAEIAYVCNIMTQRGETEHFTSDHVEVLHRLGRPFIDTVLVNIEK  
VPQEYMN5NRFDEYLVQVEHDFVGLCKQVSRVISSNFLRLENGGAFFHDGDLIVDELMRI  
IQVKKZ

(SEQ. ID. NO. 142)

MKNLIKLLIIRLIVNLADSVFYIVALWHVSNNYSSSMFLGIFIAVNYLPDLLLLIFFGPVIDRVNPQKILIISILVQLAVAVIFLLLLNQISF  
WVIMSLVFISVM5ASSISYVIEDVLIPQVVEYDKIVFANSLFSISYKVLD5IFNSFASFLQVAVGFILLVKIDIGIFLLALFILLLLKFRTSN  
ANENFSFKYKREVLQGTGKFI5LNKLLFKTSISLTINFFYSFQTVVPIFSIRYFDGPIFYGIFLTIAGLGGILGNMLAPIVIKYKLSN  
QIVGVFLFLNGSSWLVAIVIKDYTL5LILFFVCFMSKGVFNII5FNSLYQQIPPHQLLGRVNTTID5IISFGMPIGSLVAGTLIDLNIELVLI  
AISIPYFLFSYIFYTDNGLKEFSIYZ

(SEQ. ID. NO. 143)

MMSNKNKEILIFAILYTVLFMFDGVKLLASLMP5AIAANYLVYVVLALYGSFLFKDRLIQQWKEIRKTKRKFFFGVLTGWLFLILMT  
VVFEFVSEMLKQFVGLDQGCLNQ5SNQSTFQEQLLIAVFAACVIGPLVEELFFRQVLLHYLQERLSGLLSIILVGLVFALTHMHSLAL  
SEWIGAVGYLGGGLAF5IYYVKEKENIYYPLL5HMLSNSLSLIIAISIVKZ

(SEQ. ID. NO. 144)

LKKPIIEFKNV5SKVFEDSNTKVLKDINFELEEGKFYTL5GASGSGKSTILNIIAGLLDATTGDIMLDGVRINDIPTNKR5DVTVFQSYA  
LFFHMNVFENVAFPLRLRKIDKKEIEQRVAEVLK5MVQLEGYEKRSIRKLSGGQRQ5RVAIARAI5NQPRVVLLDEPLSALDLK5LRTD  
MQYELRELQ5RLGITFV5VTHDQEEALAMSDWIFVMNDGEIVQSGTPVDIYDEPINHFVATFIGESNILPGTMIEDYLV5FENGKRFE  
AVDGGMKPN5EPVEVVIRPEDLRITLPEEGKLQV5KVD5TLFRGVHYEIIAYDELGNEWMIH5STRKAIVGEEIGLDFEPEDIHIMRLNE  
TEEEFDARIEEYVEIEEQEAGLINAIEEERDEENKLZ

(SEQ. ID. NO. 145)

MKSMRILFLLALIQLSSCFLWKECIL5SFKQSTAFFIGSMVFVSGICAGVNYLYTRKQEVH5SVLASKK5SVKLFYSMLLLINLLGAVL  
VL5DNLF5IKNTLQ5QLVDFLLPSFFL5FLD5LLIFLPLK5YVRD5FLAMLDRK5KTVLV5TILATLLFLRNPM5TIVSL5LIYIGLGLFFAA5YL  
VPNSV5KKEVS5FYGHIFRDLVLVIVTLIFFZ

(SEQ. ID. NO. 146)

MVKKIIGMVLALLSVTVVGVGVFAYTIYQ5QGTETLAKTYKKIGEETKVIEATEPLTILLMGVDTGNVERTETVWGRSD5SMILMTV  
NPKTKKTTMMSLERDILTRIESGNGQAHEAKLNSAYADGGAELAIETIQKMMNIHIDRYVMVNM5RGLQKLVD5AVGGITVNNILGF  
PISISDQEEFNTISIGVGEQHIGGEEALVYARMRYQDPEG5YGRQKRQREVIQK5VMEKALSLSNSIGHYQEILKALS5DNMQTNIDLSA  
KSIPNLLGYKDSFKTIETQQLQ5GE5ILQGV5SYQIVSR5AHM5LEMQNLLRRSLGQEEVTQLETNAVL5FEDL5FGRAPV5GDEDNZ

(SEQ. ID. NO. 147)

MKKQAYV5IIALTSFLVFFFSHSLLEILD5FDWSIFLHDVEKTEKFV5L5LVFSMSMTCLLALFWRGIEEL5LRKM5QANLKRLLAGQE  
VVQVADPDLDASFK5LSGK5LNL5TEALQKAENQSLAQEE5IIEKERKRIARDLHDTVSQELFAAHMIL5SGISQ5QAL5KL5DREKM5QTQ  
LQSVTAILETAQKDLRVLLHLRPVELEQKSLIEGIQILLKELEDK5DLRVSLKQNM5TKLPK5KIEEHIFRILQELISN5TLRHAQASCLD  
VYLYQTDV5ELQLKVVDNGIGFQLGSLDDLSYGLRNIKERVEDMAGTVQLLTAPKQGLAVDIRIPLLDKEZ

(SEQ. ID. NO. 148)

MIVSII5QG5FVWAILGLGIFMTFRILNFPDMTTEGSFPLGGAVAVTLITKGVNPFLATLVAVGAGCLAGMAAGLLYTKGKIPTLLSGI  
LVMTSCHSIMLLIMGRANLGLL5GKQIQDVL5PFD5DLNQLLTGLIFVSIVIALMLFFLDTKL5GQAYIATGDNPDMARS5FGIHTGRME  
LMGLVLSNGVIALAGALIAQ5QEGYADVSRGIGVIVVGLASLIIGEVIFK5LSLAERLVTIVVGSIA5YQFLVWAVIALGFNTSYLRLYS  
ALILAVCLMIPTFKQ5TILKGAKLSKZ

(SEQ. ID. NO. 149)

MKKMKVWSTVLATGVALTTLAACSG5SNSTTAS5SEEKADK5QELVIYSNSV5SNGRGDWLTAKAKEAGFNIK5MVDIAGAQ5LADR  
VIAEKNN5AVADMVFGIGAVDSNKIRDQKLLVQYKPKWLDKIDQSL5DKDN5YNPVIVQPLVLIGAPDV5KEMPKDWTEL5GSKYK5G  
KYSISGLQGGTGRAILASILVRYLDDKGELGVSEKGWEV5AKEYLK5NAYTLQK5ESSIVKMLDKEDPIQYGMW5G5GALV5GQKEQ  
NVVFKVMTPEIGVPFVTEQTMVL5STSKQALAKEFIDWFGQ5SEIQVEY5SKNFGSIPANKDALKDLPEDTKKFVDQV5KPNIDWEA  
VGKHLDEWVEKAELEYVQZ

(SEQ. ID. NO. 150)

MIKFDNIQIKY5GDFVAIDN5LNLDIHEGEFFTLG5PSGCGK5STL5RALV5GLDPSSGSIEVNGT5DVT5HLEPEKRGIGIVFQ5SYALFPTMT  
VFDNIAFGLKV5KKVAPDV5IKAVSAVAAKISDQQLQRNV5ELSGGQ5QRVALARALVLEPKILCLDEPL5NLDAKL5RVDLR5KEL  
KRLQKELGITTLYVTHDQ5EAL5L5DRIA5VF5N5GYIEQV5GTPVEIYHNSQTEFVCDFIGDINVLTD5TVHEVLLKNT5VFLEDK5GYI  
RLEKVRFNRETEQDFILKGTIIDV5FSGV5TIHYTIKVS5ESQILN5VTSID5QA5IRS5VGESVELFITPSDVLQFZ



(SEQ. ID. NO. 151)

MRHKLNLDWLIRLGLIWFLVTFHIYPNFDLVVNVFVKGGFSLDAVHRVLKSQRALQSIMNSFKLAFSLIITVNVVGILCVLFTEYF  
DIKGAKILKGYMTSLIYGGVVLATGYKFVYGPYGLITKFLQNVIPSLDPNWFIFYGAVLFIMTFSGTANHTLFLTNTIRSVDYHTIE  
AARNMGAKPFTVFRKVVLPTLIPTLFALTIMVFLSGLSAVAAPMIVGGKEFQTINPMIITFAGMGNSRDLAALLAILGIATTILLTIM  
NKIEKGGNYISISKTAPLKKQKASKPWNIIAHIVAYGLFTVFMLPLIFIVLYSFTDPVAIQTGNTLSNFTLENYRLFFSNSAAFSPF  
LVSEIYSIIAATTATILAVVFARVVRKHKSDFDFEYGALLPWLLPSTLLAVSLLFTFNQPQFLVLNQILVGSVLILLIAIYIVVKIPFSY  
RMVRAILFSVDDMEDAARSMGASPFYTMMKVHIFILPVVLSVIALNFNSLLTDFDLSVFLYHPLAQPLGITIRSAGDETATSNAQA  
LVFVYTIVLMIISGTVLYFTQRPGRKVRKZ

Table 3

**ID201 - 4106.4 (SEQ. ID. NO. 168)**

ATGATAAAAAATCCTAAATTATTAACCAAGTCTTTTTTAAAGAGTTTTCGAATTCCTAGGTGGTGGTGGTCTAGTCATTCATAT  
 AGCTATTTATTTGACCTTTCTTTTATTATATTCAACTGGAGGGGAAAAAGTTTAAATGAGAGCGCAAGAGTGTTTACGGAGT  
 ATTTAAAGACTAAGACATCTGATGAAATTCCAAGCTTACTCCAGTCTTATTCAAAGTCCTTGACCATATCTGCTCACCTTAA  
 AGAGATATTGTAGATAAGCGGCTCCCTCTTGTCATGACTTGGATATTAAAGATGGAAAGCTATCAAATTATATCGTGATGTT  
 AGATATGTCTGTAGTACAGAGATGGTAAACAGGTAACCGTGCAATTTGTTTACGGGGTGGATGTCTACAAAGAAGCAAAGA  
 ATATTTTGTCTTTTGTATCTCCATATACATTTTGGTTACAATTGCTTTTTCTTTGTTTCTTATTTTATACTAAACGC  
 TTGCTCAATCCTCTTTTTTACATTTTCAAGGTGACTAGTAAATGCAAGATTGGATGACAATATTCGTTTGTATGAAAGTAG  
 GAAAGATGAAGTTGGTGAAGTTGAAAAACAGATTAAATGGTATGTATGAGCACTTGTGAAGGTTATTTATGAGTTGGAAGTC  
 GTAATGAGCAAATTGTAAATTTGCAAAATCAAAAGGTTTCTTTGTCCGCGGAGCATCACATGAGTTGAAAACCCCTTTAGCC  
 AGTCTTAGAATTATCCTAGAGAATATGCAGCATAATATTGGAGATTACAAAGATCATCCAAATATATTGCAAGAGTATAAA  
 TAAGATTGACCAGATGAGCCACTTATTAGAAGAAGTACTGGAGTCTTCTAAATCCAAAGAGTGGACAGAGTGTGCTGAGACCT  
 TGACTGTTAAGCCAGTTTGTAGTAGATATTTTATCACGTTATCAAGAATTAGCTCATCAATAGGTGTTACAATTGAAAATCAA  
 TTGACAGATGCTACCAGGTCGTCATGAGTCTTAGGGCATTTGGATAAGGTTTTGACAAAACCTGATTAGTAATGCAATTAAATA  
 TTCAGATAAAAAATGGGCGTGTAAATCATATCCGAGCAAGATGGCTATCTCTCTATCAAAAATACATGTGCGCCTCTAAGTGACC  
 AAGAACTAGAACATTTATTGATATATTCTATCTCAATCGTGACAGATAAGGATGAAAGTCCGGTTTGGGTCTTTAC  
 ATTGTGAATAATATTTTAGAAAGCTATCAAATGGATTATAGTTTCTCCCTTATGAACACGGTATGGAATTAAGATTAGCTT  
 GTAG

**(SEQ. ID. NO. 152)**

MIKNPKLLTKSFLRSFAILGGVGLVIHIAIYLTFPPFYIIQLEGEKFNESARVFTEYLKTKTSDEIPSLLSYSKSLTISAHLK  
 RDIVDKRLPLVHDLIDKDGKLSNYIVMLDMSVSTADGKQVTVQFVHGVVDVYKEAKNILLLYLPYTFVLTIAFSFVFSYFYTKR  
 LLNPLFYISEVTSKMQDLDDNIRFDESKRDEVGEVKGINGMYEHLKVIYELESNEQIVKLQNKVSFVRGASHELKTPLA  
 SLRIILENMQHNIQDYKDPKYIAKSINKIDQMSHLLLEEVLESSKFQEWTECRETLTVKPVLDVILSRVQELAHSIGVTIENQ  
 LTDATRVVMSLRALDKVLTNLISNAIKYSKNGRVIISEQDGYLSIKNTCAPLSDQLEHLFDIFYHSQIVTDKDESSGLGLY  
 IVNNILESYQMDYSFLPYEHGMEFKISLZ

**ID202 - 4106.9 (SEQ. ID. NO. 169)**

ATGGATAAAATTATTTAAACTATATCAGAAAGCGGAGCCTTTCTGCTTTTGTCTTGTATAGCACTGAAACCGTCCGCACTGC  
 TCAAGAAAAACATCAAACCAAGCTAGCTCAACTGTAGCGCTTGGTTCGAACCTTATCGCTAGCCAGATTCTCGCAGCCAAATG  
 AAAAAGGAAATACCAAACCTTACAGTTAAGGTGTTGGGATCTAGCTCTCTAGGTGCTATTATCACCGTCGCTGATACCAAGGGG  
 AACGTCAAAGGCTATGTTCAAATCCTGGTGTGACATCAAAAAGACTGCGACTGGTGAAGTCCTAGTCGGACCTTTTGTGTTG  
 AAATGGTCAATTCCTCGTTATCACAGACTACGGTACTGGAAATCCTTACAACCTATAACTCCCCTCATCTCTGGAGAAATCG  
 GTGAAGACCTTGCCCTTTTACCTTACTGAAAGCCAACAAACGCTTCAAGCGTCCGCTCAATGTCTTTTGGACGAGGAAGAC  
 AAGGTCAAGGTGTCAGGTGGTTCCTAGTTCAAGTCTTCCAGGAGCCAAGAAAGAGAGATTGCTCGCTTTGAAAAACCGCAT  
 CCAAGAAATGCCAGCTATCTCTACTCTTCTCGAAAGCGACGACCATATCGAAGCCCTCCTCAAGGCTATCTACGGGGACGAAG  
 CCTACAAGCGTCTTTCTGAAGAAGAAATCCGTTTCCAATGTGACTGTAGCCATGAACGCTTTATGAACGCTCTTGCCAGCCTT  
 CCAAGCTCAGACTTACAGGAAATGAAAGAGGAAGACCACGGGGCAGAAATCACTTGTCAATTCGCCAAACTACTTACAACCTT  
 TGATGAAAAGGACCTGGAGGAACCTATTCTGTGACAAATCTTAA

**(SEQ. ID. NO. 153)**

MDKIKTISEGAFRAFLVDSTETVRTAQEKHQQTASSTVALGRTLIAISQILAAANEKGNKLTIVKVLGSSSLGAIITVADTKG  
 NVKGYVQNPQVDIKKTATGEVLVGPFGVNGQFLVITDYGTPNPYNSITPLISGEIGEDLAFYLTESQTTSAVGLNVLLDEED  
 KVKVAGGFLVQVLPQAKKEEIAFEKRIQEMPAISTLLESDDHIEALLKAIYGD EAYKRLSEEEIRFQCDCSHERFMNALASL  
 PSSDLQEMKEEDHGAEITCQFCQTTYNFDEKDLLEELIRDKSZ

**ID203 - 4115 (SEQ. ID. NO. 170)**

ATGAAATCAATAACTAAAAAGATTAAAGCAACTCTTGCAAGAGTAGCTGCCTTGTGTTGCAGTATTTGCTCCATCATTTGTATC  
 TGCTCAAGAATCATCAACTTACACTGTAAAGAAGGTGATACACTTTTCAAGAAATCGCTGAAACTCACAACACAACAGTTGAAA  
 AATTGGCAGAAAACAACCAATTGATAACATTCAATTGATTTATGTTGATCAAGAGTTGGTTATCGATGGCCCTGTAGCGCCT  
 GTTGCAACACCAGCGCCAGCTACTTATGCGGCACCAGCCGCTCAAGATGAAACTGTTTCAAGCTCAGTAGCAGAAATCCAGT  
 AGTAAGTGAACAGTTGTTTCAACTGTAAGCGGATCTGAAGCAGAAGCCAAAGATGGATCGCTCAAAAAGAATCAGGTGGTA  
 GTATACAGCTACAAATGGACGTTATATCGGACGTTACCAATTAA

**(SEQ. ID. NO. 154)**

MKSITKKIKATLAGVAALFAVFAPSFVSAQESSTYTVKEGDTLSEIAETHNTTVEKLAENNHIDNIHLIYVDQELVIDGPVAP  
 VATPAPATYAAPAAQDETVSAPVAETPVVSETVSVSGSEAEAKEWIAQKESGGSIQQLQMDVISDVNTZ

**ID204 - 4117.1 (SEQ. ID. NO. 171)**

ATGAATTTAGGAGAATTTTGGTACAATAAAATAAATAAGAACAGAGGAAGAAGGTTAATGAAGAAAAGTAAGATTTATTTT  
 AGCTCTGCTATTTTCTTAGCTAGTCCAGAGGGTGCAATGGCTAGTGATGGTACTTGGCAAGGAAAAACAGTATCTGAAAGAAG  
 ATGGCAGTCAAGCAGCAAAATGAGTGGGTTTTGATACTCATTATCAATCTTGGTTCTATATAAAGCAGATGCTAACTATGCT  
 GAAAATGAATGCTTAAAGCAAGGTGACGACTATTTTACCTCAAATCTGGTGGCTATATGGCCAAATCAGAATTGGGTAGAAGA  
 CAAGGGAGCCTTTTATTATCTTGACCAAGATGGAAGATGAAAAGAAATGCTTGGGTAGGAACTTCCATGTTGGTGAACAG  
 GTGCCAAAGTAATAGAAGACTGGGTCTATGATTCTCAATACGATGCTTGGTTTATATCAAAGCAGATGGACAGCAGCAGAG  
 AAAGAATGGCTCCAAATTAAGGGAAGGACTATTATTTCAAATCCGGTGGTTATCTACTGACAAGTCAGTGGATTATCAAGC  
 TTATGTGAATGCTAGTGGTGCCAAAGTACAGCAAGGTGGCTTTTTTGACAAACAATACCAATCTTGGTTTATCATCAAAGAAA  
 ATGGAACTATGCTGATAAAGAAATGGATTTTCGAGAATGGTCACTATTATATCTAAATCCGGTGGCTACATGGCAGCCAAT  
 GAATGGATTTGGGATAAGGAATCTTGGTTTTATCTCAAATTTGATGGGAAAAATGGCTGAAAAGAAATGGGTCTACGATTCTCA  
 TAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTACATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTATC

TCAAATCTGATGGGAAAATAGCTGAAAAAGAATGGGCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGT  
TACATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTTACCTCAAATCTGATGGGAAAATAGCTGAAAAAGAATG  
GGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCTGGTGGCTACATGGCGAAAAATGAGACAGTAGATGGTTATC  
AGCTTGGAGCGATGGTAAATGGCTTGGAGGAAAAACTCAAATGAAAATGCTGCTTACTATCAAGTAGTGCTGTACAGCC  
AATGTTTTATGATTAGATGGTGAAGAGCTTTCTATATATCGCAAGGTAGTGTGCTATGGCTAGATAAGGATAGAAAAAGTGA  
TGACAAGCGCTTGGCTATTACTATTTCTGGTTTGTACGGTATGAAAAACAGAAGATTACAAGCGCTAGATGCTAGTAAGG  
ACTTTATCCCTTATTATGAGAGTGATGGCCACCGTTTTTATCACTATGTGGCTCAGAATGCTAGTATCCCAGTAGCTTCTCAT  
CTTTCTGATATGGAAGTAGGCAAGAAATATTATTCGGCAGATGGCCTGCATTTTGTATGGTTTAAAGCTTGAGAATCCCTTCTT  
TTTCAAAGATTTAACAGAGGCTACAAACTACAGTGCTGAAGAATTGGATAAGGTATTTAGTTTGCTAAACATTAACAATAGCC  
TTTTGGAGAACAAAGGGCGCTACTTTTAAGGAAGCCGAAGAACATTACCATATCAATGCTCTTTATCTCTTGGCCATAGTGCC  
CTAGAAAGTAACTGGGGAAGAAGTAAATTGCCAAAGATAAGAATAATTTCTTTGGCATTACAGCCTATGATACGACCCCTTA  
CCTTTCTGCTAAGACATTTGATGATGTGGATAAGGGAATTTTAGGTGCAACCAAGTGGATTAAGGAAAATTATATCGATAGGG  
GAAGAACTTTCTTGGAAACAAGGCTTCTGGTATGAATGTGGAATATGCTTCAGACCCTTATTGGGGCGAAAAAATGCTAGT  
GTGATGATGAAAATCAATGAGAAGCTAGGTGGCAAAGATTAG

**(SEQ. ID. NO. 155)**

MNLGEFWYNKINKNRGRRLMKKVRFI FLALLFFLASPEGAMASDGTWQKQYLKEDGSQAANEWVFDTHYQSWFY IKADANYA  
ENEWLKQDDYFYFKSGGYMAKSEWVEDKGAFYYLDQDGKMKRNVWGTSYVGATGAKVIEDWVYDSQYDAWFI IKADQHA  
KEWLQIKGKDYFFKSGGYLLTSQWINQAYVNASGAKVQGGWLDKQYQSWFY IKENGNVADKEWIFENGHYIYLLKSGGYMAAN  
EWIWDKESWFIYLFKDGKMAEKWVYDSHSQAWYFYSKSGGYMTANEWIWDKESWFIYLSKDGKIAEKWVYDSHSQAWYFYSKSGG  
YMTANEWIWDKESWFIYLSKDGKIAEKWVYDSHSQAWYFYSKSGGYMAKNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTA  
NVYDSGEKLSYISQGSVWVLDKDRKSDDKRLAITISGLSGYMKTEDLQALDASKDFIPYIESDGHFRFYHYVAQNASIPVASH  
LSDMEGKKYYSDFGLFDPGFLENPFLLKDLTEATNYSAEELDKVPSLLNNINSLLENKGATFKEAEEHYHINALYLLAHS  
LESNWGRSKIADKNNFPGITAYDTTPYLSAKTFDDVDKIGILGATKWIKENYIDRGRFTLGNKASGMNVEYASDPYWGKIAS  
VMMKINEKLGGKDZ

**ID205 - 4118.1 (SEQ. ID. NO. 172)**

ATGAAAAAATAGGTACATTACTCGTTCTCTTCTTTCTGCAATCATTCTTGTAGCATGTGCTAGCGGAAAAAAGATACAAC  
TTCTGGTCAAAAACATAAAGTTGTTGCTACAACTCAATCATCGCTGATATTACTAAAAATATTGCTGGTGACAAAATGACC  
TTCATAGTATCGTTCCGATTGGGCAAGACCCACACGAATACGAACCACTTCTGAGACGTTAAGAAAACTTCTGAGGCTAAT  
TTGATTTTCTATAACGCTATCAACCTTGAACAGGTGGCAATGCTTGGTTTACAAAATTTGGTAGAAAATGCCAAGAAAACCTGA  
AAACAAGACTACTTCGAGTCAGCGACGGCGTGTGATGTTATCTACCTTGAAGGTCAAATGAAAAAGGAAAAAGAACCCAC  
ACGCTTGGCTTAACTTGAACACGGTATTATTTTGTCTAAAAATATCGCCAAACAATTGAGCGCCAAAGACCTTAAACAATAA  
GAATTCATGAAAAAATCTCAAAGAATATACTGATAAGTTAGACAACTTGATAAAGAAAGTAAGGATAAATTTAATAAGAT  
CCCTGCTGAAAAAGAACTCATTTGTAACAGCGAAGGAGCATCAAATCTTCTTAAAGCCTATGGTGTCCCAAGTGCTTACA  
TCTGGGAAATCAATGATGAAGAAGGAACCTCTGAACAAATCAAGACCTTGGTTGAAAAACTTGCCCAACAAAGTTCCA  
TCACTCTTTGTAGAAATCAAGTGTGGATGACCGTCCAATGAAAACTGTTTCTCAAGACACAAACCTCCCAATCAAGTCAAT  
CTTTACTGACTCTATCGCAGAACAAGGTAAAGAAGGCAGCTACTACAGCATGATGAAATACAACCTTGACAAGATTGCTG  
AAGGATTGGCAAAATAA

**(SEQ. ID. NO. 156)**

MKKLGTLLVFLSAIILVACASGKKDTSQGLKVVATNSIIADITKNIAGDKIDLHSIVPIGQDPHEYEPLPEDVKKTSEAN  
LIFYNGINLETGGNAWFTKLVENAKTKNDYFAVSDGVDVILEGQNEKGKEDPHAWLNLENGIIFAKNIQKLSAKDPNNK  
EFYEKLNKEYTDKLDKDKESKDKFNKI PAEKKLIVTSEGAFFYFSKAYGVPSAYIWEINTEEEGTPEQIKTLVEKLRQTKVP  
SLFVESSVDDRPMTVSQDTNIPYIQAIFDTSIAEQGKEGDSYYSMMKYNLDKIAEGLAKZ

**ID206 - 4119.1 (SEQ. ID. NO. 173)**

ATGGAATGGTATAAAAAAATCGGACTTCTTGCAACTACAGGTTTAGCTTTGTTTGGGCTCGGCGCTTGCTCCAACATATGGTAA  
ATCTGCGGATGGCACAGTACCATCGAGTATTTCAACCAGAAAAAGAAATGACCAAAACCTTGAAGAAATCACTCGTGATT  
TTGAGAAGGAAAAACCTTAAGATCAAGGTCAAAGTCGTCAATGTACCAATGTGCTGGAAGTATTGAAGACACGCGTTCTCGCA  
GGAGATGTGCCTGATGTGGTCAATATTTACCCACAGTCCATCGAAGTCAAGAAATGGGCAAAAGCAGGTGTTTTTGAAGATTT  
GAGCAACAAAGACTACCTGAAACGCGTGAAAAATGGCTACGCTGAAAAATATGCTGTAAACGAAAAAGTTTACAACGTTCTCTT  
TTACAGCTAATGCTTATGGAATTTACTACAACAAAGATAAATTCGAAGAAGTGGGCTTGAAGGTTCTGAAACCTGGGATGAA  
TTTGAACAGTTAGTCAAAGATATCGTTGCTAAAGGACAAACACCATTTGGAATTCAGGTGTCAGATGCTTGGACACTCAATGG  
TTACAATCAATTAGCTTTTGCAGACAGCAACAGGTGGAGGAAAAAGCAAAATCAATACCTTCGTTATTTCTCAACCAATGCCA  
TTAAATTTGTCGGATTCGATTATGAAAGATGATATCAAGGTCAAGGATCATGACATCTTCGCATCAATGGATCTAAGCAAAAGAACTGG  
GAAGGTGCTGGCTATACCGATGTTATCGGAGCCTTCGCACGTGGGGATGTCCTCATGACACCAATGGGTCTTGGGCGATCAC  
AGCGATTAATGAACAAAAACCGAATTTAAGATTGGGACCTTCATGATTCCAGGAAAAAGAAAAAGGACAAAGCTTAACCGTTG  
GTGCGGAGACTTGGCATGGTCTATCTCAGCCACCACCAAAATCCAAAAGAGCCAATGCCTTTGTGGAATATATGACCCGT  
CCAGAAGTCATGCAAAAAATACTACGATGTGGACGGATCTCCAACAGCGATCGAAGGGGTCAAACAAGCAGGAGAAGATTACCC  
GCTTGCTGGTATGACCGAATATGCCTTTACGGATCGTCACTTGGTCTGGTTGCAACAATACTGGACCAGTGAAGCAGACTTCC  
ATACCTTGACCATGAACATATGTCTTGACCGGTGATAAAACAAGGCATGGTCAATGATTGAAATGCCTTCTTTAACCCGATGAAA  
CGGATGTGGATTAG

**(SEQ. ID. NO. 157)**

MEWYKIGLLATTGLALFGLGACSNYGKSADGVTITIEYFNQKEMTKTLEEITRDFEKENPKIKVKVNVNPNAGEVLKTRVLA  
GDVPDVVNIYPQSIELQEWAKAGVFEDLSNKDYLKRVKNGYAEKYAVNEKVVNVPFTANAYGIYNNKDFEELGLKVPETWDE  
FEQLVKDIAVAKGQTFPGIAGADAWTLNGYNQLAFATATGGGKEANQYLYRYSQPNAIKLSDPIMKDDIKVMDILRINGSKQKNW  
EGAGYTDVIGAFARGDVLMTPNGSWAITAINEQKPNFKIGTFMIPGKEKGQSLTVGAGDLAWSISATTKHPKEANAFVEYMT  
PEVMQKYDVGDSPTAIEGVKQAGEDSPLAGMTEYAFTRHLVWLQYWTSEADFHTLTMNYVLTGDKQGMVNDLNAFFNPMK  
ADVDZ

**ID207 - 4123.1 (SEQ. ID. NO. 174)**

ATGAAGAAAATCAAACCGCATGGACCGTTACCAAGTCAGACTCAGCTAGCTTATCTGGGAGATGAACTAGCAGCTTTTATCCA  
CTTCGGTCTCTAATACCTTTTATGACCAAGAATGGGGGACTGGACAGGAGGATCCTGAGCGCTTTAACCCGAGTCAGTTGGATG  
CGCGTGAGTGGGTTCTGTGCTCAAGGAAACGGGCTTCAAAAAGTTGATTTTGGTGGTCAAGCACCACGATGGCTTTGTCTT  
TATCCGACAGCTCACACAGATTATTCGGTTAAGTCAGTCTTGGAGGAGAGGAAAGGGCGACTTGCTCCTTGAAGTATCCCA

AGCTGCCACAGAGTTTGATATGGATATGGGGGTCTACCTGTCCACCGTGGGATGCCCATAGTCCCCCTCTATCATGTGGACCGAG  
 AAGCGGACTACAATGCCTATTATCTGGCTCAGTTGAAGGAAATCTTATCAAATCCTAACTATGGGAATGCTGGTAAGTTTCGCT  
 GAGGTTTGGATGGATGGTGCCAGAGGAGAGGGCGCGCAAAGGTTAATTATGAATTTGAAAAATGGTTTGAACCATTTCGTGA  
 CCTGCAAGGGCGATTGCTTGATTTTTTCAACAGAAAGCACCAGTATCCGCTGGATTGGCAATGAACGAGGGTATGCAGGTGATC  
 CACTGTGGCAAAGGTGAATCCTGATAAACTAGGAACAGAGAGCTGAACCTATCTTCAGCACGGGGATCCCTCGGGCACG  
 ATTTTTTCAATCGAGAGGCAGATGTTTCCATCCGTCCAGGCTGGTTCTACCATGAGGATCAGGATCCTAAGTCTCTCGAGGA  
 GTTGGTCGAAATCTACTTTCTCAGTAGGGCGAGGAACCTCACTCTTGCTTAATATTCGCCCGAATCAAGCTGGGCTCTTTG  
 ATGCAAAGGATATTGAACGACTTTATGAATTTGCGACCTATCGCAATGAGCTCTATAAAGAAGATTGGCTCTGGGAGCTGAG  
 GTATCTGGTCCAGCTCTTTCCGAGACTTTGCTTGTGCGCCATTGACAGACGGCCTTGAGACCAGCTCTTGGGCAAGCGATGC  
 AGACTTGCCCATCCAGTTAGAACTCGACTTAGGTTCTCTAAACCTTTTGATGTAATTGAGTTAAGAGAAGATTGAAGCTAG  
 GGCAACGAATCGCTGCTTTTCATGTGCAAGTAGAGGTGGATGGTGTCTGGCAGGAGTTTGGTTCCGGGTCTACTGTTGGTTAC  
 AAACGCTCTTACGAGGAGCAGTTGTTGAGGCACAGAAGATACGTGTAGTATTACAGAATCACAGGCTTTGCCCTTGTGAC  
 CAAGATTTCCCTTTATAAACTCCTGGATTATCAAAAAAGAAGTTGTTTCAAGAACTAGCATTTCAGAAAAAGCCTAGCTG  
 TGGCAAAGGAGAAAAATGCCTATTTTACAGTTAAGCGCAGAGAATGTAGTGGTCTTTAGAAGCTAAGATTTCCGATTCAACCG  
 GGGACAGGTGTCCATGGTGTGCGCTATCAGGATGAGATTCAAGTCTTGCCTTTCAAACCTGGTGAGACTGAAAAAGTCTGAC  
 GCTACCAACCTTGATTTTCGACAGGAGATAAAACCTTGGATTCTATCTGAACCTAACGGTGGATGGTCAGCTTGTGGATCAAC  
 TTCAAGTCCAAGTTTCATAA

## (SEQ. ID. NO. 158)

MKKIKPHGPLPSQTQLAYLGDELAFAIHFGPNTFYDQEWGTGQEDPERFNPSQLDAREWVRVLKETGFKKLILVVKHHDGFVL  
 YPTAHTDYSVKVSPWRRKGDLLEVSQAATEFDMGMGVYLSPWDHSPHYHVDREAYNAYYLAQLKEILSNPNYGNAGKFA  
 EVWMDGARGEGAQVNYEFKWFETIRDLQGDCLIFSTEGTSIRWIGNERGYAGDPLWQKVNPKDLGTEAELNYLQHGDPST  
 IFSIGEDVSRPGWFYHEDQDPKSLEELVEIYFHSVGRGTPLLLNPPNAGLFDKDIERYEFATYKEDLDLGAEL  
 VSGPALSDAFACRHLTDGLETSWASDADLPQLLELDLGSPTFDVIELREDLKLQRIAAFHVQVEVDGVWQEFSGHTVGY  
 KRLLRGAVVEAQKIRVVITESQALPLLTKISLYKTPGLSKKEVVQELAFAEKSLAVAKGENAYFTVKRRECSGPLEAKISIQP  
 GTGVHGVAYQDEIQVLAFTGETEKSLLTPTLYFAGDKTLDFYLNLTVDGQLVDQLQVQVSZ

## ID208 - 4125.12 (SEQ. ID. NO. 175)

ATGCTTGAAGACTGAAAGAATACATTATATGTTTGGATCAGTTTAATTTTTATGATTTTCCCATCCTGTCTGTAGTGAC  
 TGGGTGGCTTTCTGCCTGGCATTATTGATTGATATTCTATTGTAGTGGCATATTTGGGTGTTTTAACAACCTAAGAGCCAGC  
 GCCTATCTTGGCTATATTGGGCGCTCATGCTGACTTATGTAGTTGGGAATACTGCCTTTGTTGCTGTTAATTATATCTGGTTT  
 TTCTTTTCTTATCCAATCTCTTAAGTTATCATTTTCAAGCTACGTAGTTTAAAGTCTTTACATGTCTGGACTTTTCTTCTTGC  
 TCAAGTCTTGTGTGGGGCAACTGTTGATTTTTCAGAGAATCGAAGTTGAGTTTCTATTCTATCTACTTGTAAATCTTACTT  
 TTGTGCTATTAAATGACTTTTGGATTGGTTCCGATTGCTATTGTCGAGGATTTGAAAGAAGCTCAGGTCAAGCAAAATGCTCAG  
 ATAAATCTATTGCTTGTCTGAAAATGAACGTAGTCGTATCGGTGAGGATTTGCATGATAGTCTGGGACATACCTTTGCTATGCT  
 GAGTGTCAAGCAGATTGTCCTTGCAGTTATTTTTCAGATGGAGGCTTATCCACAGGTGAAAAGGAATTAAAGAAATTCACC  
 AGATCAGCAAGGATCCATGA

## (SEQ. ID. NO. 159)

MLERLKRHYMFWISLIFMIFPILSVVTGWLSAWHLLIDILFVVAYLGLVLTTKSQRLSWLYWGLMLTYVVGNTAFVAVNYIWF  
 FFFLSNLLSYHFSVRSLKSLHVWTFLLAQVLVVGQLLIFQRIEVEFLFYLLVILTFVDLMTFGLVRIRIVEDLKEAQVKQNAQ  
 INLLLAENERSRIGQDLHDSLGHTFAMLSVKTDLALQLFQMEAYPQVEKELKEIHQISKDPZ

## ID209 - 4126.3 (SEQ. ID. NO. 176)

ATGAATGATAAGTTAAAAATCTTCTTGTGTGCTAGGAGTATTTTTTCTAGCCATAACCGGTTTCTATGTTCTATTGATACGAAA  
 TGCAGGCGACAGAGATGCCTCGCAAATTTGAAAAGGCGGCAGTTAGCCAAGGAGGAAAAGCAGTGAAGAAAAACAGAAATAGTA  
 AAGACGCAGACTTGCACGAAATTTATCTAGCTGGAGGTTGTTTCTGGGGAGTGGAGGAATATTTCTCACGTGTTCCCGGGGTG  
 ACGGATGCCGTTTTCAGGCTATGCAAATGGTAGAGGAGAAACAACCAAGTACGAATTGATTAACCAACAGGTCATGCAGAAAC  
 CGTCCATGTCACTTATGATGCAAGCAAATTTCTCTCAAGGAACTCTGCTTCACTATTTCCGCATTATCAATCCAACCAGCA  
 AAAATAAACAAGGAAATGATGTGGGGACCCAGTACCGTACTGGTGTATTACACAGATGACAAGGATTTGGAAGTGATTAAAC  
 CAAGTCTTTGATGAGGTGGCTAAGAAATACGATCAACCTCTAGCAGTTGAAAAGGAAACTTGAAGAATTTGTGGTGGCTGA  
 GGATTACCATCAAGACTATCTCAAGAAAAATCCAAATGGCTACTGCCATATCAATGTTAATCAGGCGGCCTATCCTGTCTATTG  
 ATGCCAGCAAAATACCAAAACCAAGTATGAGGAATTTGAAAAGACCCCTGTACCTGAGGAGTATGCAGTTACCCAGGAAAT  
 CAAACAGAACGAGCTTTCTCAAACCGTTACTGGGATAAATTTGAATCCGGTATCTATGTGGATATAGCAACTGGGGAACCTCT  
 CTTTTTCATCAAAAGACAAATTTGAGTCTGGTGTGGCTGGCCTAGTTTACCAACCCATCAGTCCAGATGTTGTACCTACA  
 AGGAAGATAAGTCTCAATATGACGCGTATGGAAGTGGGAGCCGAGTAGGAGATTCTCACCTTGGGCAATGCTTTACCGGAT  
 GGTCCACAGGACAAGGGCGGCTTACGTTACTGTATCAATAGCCTCTCTATCCGCTTTATTCCCAAAGACCAATGGAAGAAAA  
 AGgcTACGCTTATTTACTAGATTATGTTGATTAA

## (SEQ. ID. NO. 160)

MNDKLIKIFLLLVFFLAITGFYVLLIRNAGQTDASQIEKAAVSQGGKAVKKEISKDADLHEIYLAGGCFWGVVEEYFSRVPV  
 TDAVSGYANGRGETTKYELINQTHAETVHVTYDAKQISLKEILLHYFRIINPTSKNKQGNVDGTQYRTGVYTTDDKDLVIN  
 QVFDVAKKYDQPLAVEKENLKNFVVAEDYHQDYLLKKNPNGYCHINVNQAAYPVIDASKYPKPSDEELKKTLSPEEYAVTOEN  
 QTERAFSNRYWDFKESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPISPDVVYTKEDKSYNMTRMEVRSRVGDSHLGHVFTD  
 GPQDKGGLRYCINSLSIRFIPKQMEKGYAYLLDYVDZ

## ID210 - 4127.1 (SEQ. ID. NO. 177)

ATGAAAAAGAAATGGATGTATTATGCTGCTTGTCTTCTAATGAATCTGCCGATGACAGTTCATCTGATAAAGGAGACGGCGG  
 TTCGCTAGTCGTTTATTCACCAAACTCAGAGGGCTTAATTGGAGCAACTATTCCTGCCTTTGAAGAAAAATATGGTATCAAAG  
 TAGAACTGATTCAAGCTTGGTACTGGAGAACTTTTCAAAAACTAGAGTCAGAAAAAGAAAGTTCTGTAGCTGATGTTATCTTT  
 GGTGTTCTTATACACAATATACTACCCACGGAGAACTCTTTGAAAACCTATCTTCAAAGAAAAATGATAATGTTATCAAAGA  
 ATATCAAAACACAACTGGCTACTCTACTCTTATACACTAGATGGTAGTGTTTAATCGTCAACCTCTGTTAACTAAAGGCA  
 TGAACATCGAAGGATATAACGATCTTTCAAACCTGAACTAAAGGAAAAATCGCAACTGCTGACCCAGCAAACTCTTCTAGC  
 GCCTTTGCTCAATTAACAAATATGCTACAAGCTCAAGGTGGTTACAAAGATGATAAGGCTTGGTCTTATGTAAAAGATCTTTT  
 CACACTTATTGATGGTAAAAATCGGTTCAAGTTTCTAGTGTCTATAAAGTAGTCGCTGATGGAGAAATGGCTGTTGGTCTCT  
 CTTATGAAGATCCAGCAGTTAACTCTTAAATGACGGAGCTAACATTAAAGGTAGTCTATCCAAAGAGGAACCGTCTTCTCTA  
 CCTGCTAGTGTCTATCGTTTAAAAATCTAAAAATGAGAAATGCAAGAAATTTATCGATTTTATCTCTCAAGAAAT

ACAAGATACACTTGGTACAACCACTACTAACCCTCTGTTTCGTAATAATGCTAAAACAAGCGAAAACATGAAACCAATTGACA  
AAATCAAAACACTCACTGAAGATTATGATTATGTCATCAAGAATAAATCAGATATCGTTAAGAAATACAACGAAGTCTTTACA  
GATATCCAATCTAAACAGTAA

## (SEQ. ID. NO. 161)

MKKKWMYYAACSSNESADSSSDKGDGSLVVYSPNSEGLIGATIPAFEEKYGIKVELIQAGTGELFKKLESEKEVPVADVIF  
GGSYTYTTHGELFENYTSKENDNVIKEYQNTTGYSTPYTLDSVLI VNPDLTKGMNIEGYNDLFPKELKGIATADPANSSS  
AFAQLTNMLQAQGGYKDDKAWSYVKDLFTLIDGKIGSSSSSVYKVVDGEMAVGLSYEDPAVKLLNDGANIKVVYPKEGTVFL  
PASAAIVKSKNMENAKKFIDFIISQEVQDTLGTTTTNRPVRKNAKTSNMKPIDKIKTLTEDYDYVIKNKSDIVKKYNEVFT  
DIQSKQZ

## ID211 - 4127.2 (SEQ. ID. NO. 178)

ATGAGTGAGATCAAAATTATTAACGCCAAAAAATCTACCACGATGTCCTGTTATTGAGAATTTGAACATTACAATTCCAAA  
AGGAAGTCTCTTTACCCTTCTTGGAGCTTCAGGATGTGGGAAAACGACCTTCTTCGTATGATTGCAGGTTTCAACAGTATCG  
AAGGTGGAGAATTTTACTTCGTATGATACAAAAATCAATAATATGGAACCCAGCAACGCAATATCGGGATGGTTTCCAAAAC  
TACGCTATTTTCCACATTTGACTGTCCGAGACAACGTTGCTTTTGGTCTTATGCAAAAGAAGGTTCCAAAAGAAGAAATTGAT  
TCAACAGACCAACAAGTATCTTGAACCTCATGCAAAATTGCTCAATATGCGGATCGAAAGCCCGATAAATCAGTGGTGGACAAC  
AACACGTGTCACTTGGCATGCGCTTAGCGGTTAATCCAAGTGTCTCCTCATGGACGAGCCACTTAGTAATCTGGAGGCC  
AACTTCGCTTGGATATGCGTCAAGCCATCCGAGAAATCCAACACGAAGTGGGAATTACAACCTGTTTATGTACCCACGACCA  
AGAAGAAGCCATGGCTATTTAGACCAAAATTGCTGTTATGAAAGATGGGGTGATCCAACAAATCGGCCGACCAAAAGAAGTCT  
ATCATAAACCAGCTAATGAGTTTGTGGCAACCTTTATCGGACGCACAAATATTATCCCTGCCAATCTTGAAAACCGGAGCGAC  
GGCGCTTATATCGTCTTTTTCAGATGGCTATGCCCTTCGAATGCCAGCTCTTGATCAGGTTGAGGAGCAAGCTATTATGTAAG  
CATTCGTCCTCGAAGAGTTTATCAAAGATGAATCTGGAGATATTGAAGGAAGTATTAGAGATAGCGTCTATCTTGGACTAAATA  
CGGATATTTTTCATTGAGACAGGTTTTCGCTCAAAAATTCAAGTTAGTGAAGAATCAACTTTTGAAGAAGATCTACAAAAGGC  
AATCGTATTCGTCTACGAATCAATACGCAAAAATTAAACATCTTTCTGCAGATGGTTCCTCAAAACCTGATAAAGGAGTCAA  
CCATGGAACGTAA

## (SEQ. ID. NO. 162)

MSEIKIINAKKIYHDPVPIENLNIIPKGSFLTLLGASGCGKTTLLRMIAGFNSIEGGEFYFDDTKINNMEPSKRNIQMVFN  
YAFPHLTVRDNVAFGLMQKKVPKEELIQNTNKYLELMQIAQYADRKPDKLSGGQQQRVTALCALAVNPSVLLMDEPLSNLEA  
KLRLDMRQAIQREIQHEVGITTVYVTHDQEEAMAIQSDQIAVMKDVIIQYIGRPKELYHKPANEVATFIRGTNIIPANLEKRS  
GAYIVFSDGYALRMPALDQVEQAIHVSIRPEEFIKDESQDIEGTIRDSVYLGNTDYFIETGFASKIQVSEESTFEEDLQKG  
NRIRLRINTQKLNIFADGSQNLIKGVNHTZ

## ID212 - 4136.1 (SEQ. ID. NO. 179)

ATGAAGAAAAAATTATTGGCAGGTGCCATCACACTATTATCAGTAGCAACTTTAGCAGCTTGTTTCGAAAGGGTCAGAAGGTGC  
AGACCTTATCAGCATGAAAGGGGATGTCTTACAGAACATCAATTTTATGAGCAAGTGAAAAGCAACCTTCAGCCCAACAAG  
TCTTGTTAAATATGACCATCCAAAAAGTTTGTGAAAAACAATATGGCTCAGAGCTTGATGATAAGAGGTTGATGATACTATT  
GCCGCTTATGAAAAAACAATATGGCGAAAACTACCAACGTGTCTTGTCAACAAGCAGGTATGACTCTTGAAAACCGTAAGCTCA  
AATTCGTACAAGTAAATTAGTTGAGTTGGCAGTTAAGAAGGTAGCAGAAGCTGAATTGACAGATGAAGCCTATAAGAAAGCCT  
TTGATGAGTACACTCCAGATGTAAACGGCTCAATCATCCGTCTTAATAATGAAGATAAGGCCAAAGAAGTTCTCGAAAAAGCC  
AAGGCAGAAGGTGTGATTTTGCTCAATTAGCCAAAGATAATTCAACTGATGAAAAAACAAGAAAAATGGTGGAGAAATTAC  
CTTTGATTCTGCTTCAACAGAAGTACCTGAGCAAGTCAAAAAGCCGCTTTTCGCTTATGATGTGGATGTTTCTGATGTGA  
TTACAGCAACTGGCACACAAGCCTACAGTAGCCAATATTACATTGTAAAACCTCACTAAGAAAAACAGAAAAATCATCTAATATT  
GATGACTACAAAGAAAAAATAAAACTGTTATCTTGACTCAAAAACAAAATGATTCAACATTTGTTCAAAGCATTATCGGAAA  
AGAATTGCAAGCAGCGCAATATCAAGGTTAAGGACCAAGCCTTCCAAAATATCTTTACCCAATATATCGGTGGTGGAGATTCAA  
GCTCAAGCAGTAGTACATCAAACGAATAG

## (SEQ. ID. NO. 163)

MKKKLLAGAITLLSVATLAACSKGSEGADLISMKGDVITEHQFYEQVKSNPQAQQVLLNMTIQKVFEKQYGSLEDDKEVDDTI  
AEEKQYGENYQVRLSQAGMTLETRKAQIRTSKLVELAVKKVAEAEALTDEAYKKAFDEYTPDVTAQIIRLNNEDKAKEVLEKA  
KAEGADFAQLAKDNSTDEKTKENGGEITFDSASTEVPEQVKKAALFDVDGVSVDVITATGTQAYSSQYIVKLTKKTEKSSNI  
DDYKEKLKTIVILTQKQNDSTFVQSIIGKELQANIKVKDQAFQNIPTQYIGGGDSSSSSSTSNEZ

## ID213 - 4137.3 (SEQ. ID. NO. 180)

ATGAAAAAATATTAACAATATGTAAACCTTAGGTACTGTAGTGGTATTATCAGCATTTGTTGCTAACTCAGTTGCAGCTCA  
GGAGACTGAAACTTCTGAAGTATCAACACCAAGTTGGTGCAACCTGTTGCACCAACGACTCCGATTCGGAAGTACAACCTA  
CATCGGATAACTCTCGGAAGTTACTGTACAACCTCGAACAGTTGAAACTACTGTTAAGGATCCATCTCTACAGCGGAAGAA  
ACTCCTGTCTTAGAAAAAATAATGTTACTTTAACAGGGGGCGGAGAAAAATGTTACTAAAGAGTTAAAGGATAAATTTACTAG  
CGGTGACTTTACTGTAGTGATTAGTACAATCAGTCAAGTGAGAAAGGCTTACAAGCTCTGTTTGGAAATATCTAATTTCAAAC  
CCGCTCAACAAAATAGTTATGTAGATGTGTTCTTAGAGACAATGGTGAGTTGGGGATGGAAGCGCGTGATACTTCTTCCAAT  
AAAAATAACCTAGTATCCAGACCTGCTTCAGTTTGGGGTAAAGTACAAAACAAGAGGCTGTGACTAACACTGTTGTCAGTAGTAGC  
AGATTCAAGTCAAAAAACAATATTTTATACGCAATGGTACAAAAGTAGTAGAAAAGAAAGTGGATATTTTCTTAACATCA  
AGGATATTAAAGGTATTGATTACTATATGCTTGGGGGAGTGAAACGTGCAGGAAAAACCGCGTTTGGTTTTAACGGAACACTA  
GAAAATATCAAAATCTTTAATAGTGCAATTGGATGAAGAACTGTTAAAAAGATGACAAACAAACGCTGTACTGGACATTTAAT  
TTATACGGCTAATGATACAACAGGTTCTAATACTTTCCGTATTCCAGTTCTGTATACTTTTAGCAATGGTTCGGGTATTTTCAA  
GCATTGACGCTCGTTACGGTGGAACTCATGATTTCTGAATAAAATTAATATTGCTACAAGTTATAGTGATGATAATGGTAAG  
ACATGGACTAAACCAAAATTAACATTGGCATTGATGATTTTGCGCCAGTACCATTAGAATGGCCTCGTGAAGTTGGTGGACG  
TGACTTACAAATCAGCGGTGGTGCAACCTATATTGACTCTGTTATTGTTGAAAAAAGAACAACAAGTACTCATGTTTGTCTG  
ATGTTAGCCTGCTGGAGTAAGTTTATAGAGAAGCAACTAGAAAAGATTCAAGTTATAAAACAAATTGATGGTAATTATTACCTT  
AAATTAAGGAACAAGGTGATCTGATTACAATTATACTATTCTGTGAGAATGGTACTGTATACGACGATCGTACCAACAGACC  
AATGAATTTTTCAGTAGATAAAAAATTTCCGGTATTAAACAAAATGGTAATTATTGACGGTAGAGCGG

## (SEQ. ID. NO. 164)

MKKNIKQYVTLGTVVVLSAFVANSVAAQETETSEVSTPKLVQVPAPTTPISEVQPTSDNSSEVTVQPRVETTVKDPSSSTAEETPVLEKNNVTLTGGENVTKELKDKFTSGDFTVVIKYNQSSSEKGLQALFGISNSKPGQQNSYVDVFLRDNGELGMEARDTSSNKNNLVSRPASVWVGKYKQEAVENTVAVVADSVKKTYSLYANGTKVVEKKVDNFLNFKDICKGIDYYMLGGVVKRAGKTAFGFNGTLENIKFFNSALDEETVKMMTTNAVTHGLIYTANDTTGSNYFRIPVLYTFSNGRVFSSIDARYGGTHDFLNKINIATSYSDDNGKWTWKPKLTLAFDDFAPVPLEWPREVGGRDLOISGGATYIDSVIVEKKNQVLMFADVMPAGVVSFREARTRKDSGYKQIDGNYYLKLKQKQDIDYNYTIRENGTVYDDRTNRPTFEVSVDKNFGIKQNGNYLTVER

**ID214 - 4185 (SEQ. ID. NO. 181)**

ATGAAAAAATTTAGCCTATTACTAGCTATCCTACCATTTTTTGGTTGCTGTGAGAATCAAGCTACACCCAAAGAGACTAGCGCTCAAAGACAATCGTCCCTTGCTACAGCTGGCGACGTGCCACCATTGACTACGAAGACAAGGGCAATCTGACAGGCTTTGATATCGAAGTTTAAAGGCAGTAGATGAAAAACTCAGCGACTACGAGATTCAATTCCAAAGAACCGCTGGGAGAGCATCTTCCCAAGGACTTGATTCTGGTCACTATCAGGCTGCGGCCAATAACTTGAGTTACACAAAAGAGCGTGCTGAAAAATACCTTTACTCGTTCCAATTTCCAACAATCCCCTCGTCTTGTGAGCAACAAGAAAAATCCTTTGACTTCTCTTGACCAGATCGCTGGTAAAAACAACACAGAGGATACCGGAACCTTCAACGCTCAATTCATCAATAACTGGAATCAGAAACACACTGATAATCCCGCTACAATTAATTTTTCTGGTGAGATATTGGTAAACGAATCCTAGACCTTGCTAACGGAGAGTTTGATTCTCTAGTTTTTGACAAGGTATCCGTTCAAAGATTATCAAGGACCGTGGTTTAGACCTCTCAGTCGTTGATTTACCTTCTGCAGATAGCCCCAGCAATTATATCATTTCTCAAGCGACCAAAAAGAGTTTAAAGAGCAATTTGATAAAGCGCTCAAAGAACCTTATCAAGACGGAACCTTGAAAACTCAGCAATACCTATCTAGGTGGTTCTTACCTCCAGATCAATCTCAGTTACAATAA

**(SEQ. ID. NO. 165)**

MKKFSLLLAILPFLVACENQATPKETSQKTIIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLSDYEIQFORTAWESIPLGLDSGHYQAAANNLSYTKERAELYSLPISNNPLVLVSNKKNPLTSLDQIAGKTTQEDTGTSTNAQFINNWNQKHTDNPATINFSGEDIGKRILDLANGEFDLVDKVSQKIIKDRGLDLSVDLPSADSPSNYIIIFSSDQKEFKEQFDKALKELYQDGTLEKLSNTYLGGSYLPDQSQLQZ

**ID215 - 4211.1 (SEQ. ID. NO. 182)**

ATGAAAAAATAGTTTTATATCATATCCTCACTCTTTTTTGTGTGTCTTATTTGTCTATGCTACGGCGACGAATTTTCAAAACAGTACCAGTGCTAGGCAGGTAAAAACGGAAACCTATACCTAATACAGTAACAAATGTCCTTATGACATACGCTATAATAGTATAAGTATTTTATTAGCGGTTTTTGCTTCAGAAGTATCAGTGGTCTTGACTGGTGCAAATCGCCTATCGCTAGCTAGTGAAATGCAAGAAAGTACACGTAAATTCAGGTTACTGCTGACCTAACAGATGCCGGTGTGGAAAGATTGAAGTTCTTTGAGCATTGAAGATTTACCCAATGGGCTGACCGCTGTGGCGACTCCGCAAAAAATTACAGTCAAGATTGGTAAGAAGCTCAGAAGGATAAGGTAAAGATTGTACCAAGAGATTGACCTAGTCAAATTGATAGTCCGGTACAAATTGAAAATGTGATGGTGTGAGATAAAGAAAGTGTCTATTACGAGTGACCAAGAGACATTGGATAGAATTGATAGATTATCGCTGTTTTGCCAACTAGCGGAACGTATAACAGGTAATTACAGTGGTTTCACTACCTTTGCAGGCAATCGACCGCAATGGTGTGTCTTACCGGCAGTTATCACTCCGTTTGATACAAATATGAAGGTGACTACAAAACAGTAGCACCAGATTCAAGCACATCAAATTCAGTACAAGCAGTTTATCGGAGACATCTTCGTCAACGAAAGCAACTAGTTCAAAAACGAATTAA

**(SEQ. ID. NO. 166)**

MKKNSLYIISLFFACVLFVYATATNFQNSTSARQVKETETNTVTNVPIDIRYNSDKYFISGFASEVSVVLGTANRLSLASEMQESTRKFKVTADLTDAVGVTIEVPLSIEDLPNGLTAVATPQKITVKIGKKAQKDKVKIVPEIDPSQIDSRVQIENVMVSDKEVSITSQETLDRIDKIIAVLPTSERITGNYSGSVPLQAIDRNGVVLPAVITPFDITMKVTTKPVAPSSSTSNSSSTSSSETSSSTKATSSKTNZ

**ID216 - 4127.3 (SEQ. ID. NO. 183)**

ATGTTGATTGGCGAAGGGTATCGGACTTTCCCTGTCTGATTTATACCCAATTTATTAGCGAGGTTGGAGGAAATCTGCTTTTGCAATTATGGCGATTATCATTGCTTGGCAATTTTTCTTATCCAAAAACACATTGCAAACCGCTACAGTTTCAGCATGAATCTGCTCCATCCAATTGAGCCTAAAAAACTACAAAAGGAAAAATGGCTGCCATTTATGCAACAGTCTACGGAATTATCTTTATCTCTGTTTACCTCAAATCTACTTAATTTATACCTCTTTCTTAAAAACATCAGGTATGGTATCTGTTAAAGGTTATTCTCCAAACAGTTACAAGGTAGCTTTCCATCGTATGGGATCTGCTATTTTCAATACCATTCTGATCCCTTTGATTGCCTTAGTTCTAGTTGTTCTATTTGCGACATTTATCTCTACCTAGCCGTTAGAAAACGGAATTTGTTTACAACTTAATTGACAGCCTCAGTATGGTACCTTATATTGTACCAGGAACCGTTCTAGGGATTGCCTTCATTTCTTCTCTCAATACTGGTCTATTTGGAAGTGGATTTCTTATGATTACAGGACTGCTTTCATCTTGATTATGTCTCTATCTGCCAGAAGATTACCTTATACTATTCTGCTCATCTGTTGCTAGCTTACAACAAATAGCACCAAGTATTGAAGAAGCTGCTGAAAGCTTAGGAAGTAGTCGCTCAATACCTTTGCTAAGATTACAACCTCCAATGATGCTATCTGGTATCATTTCTGGAGCCATCTTATCTTGA

**(SEQ. ID. NO. 167)**

MLIGEGYRTPVLIYTFQFISEVGGNSAFAIMAIIIALAIFLIQKHIANRYSFSMNLHPIEPKKTTKGKMAAIYATVYGIIFI SVLPQIYLIYTSFLKTSGMVSVKGYSPNSYKVAFHRMGSAIFNTIRIPLIALVLVLPATFISYLAVRKRNLFTNLIDSLSMVPIVPGTVLGIAFISSFNTGLFGSGFLMITGTAFILIMSLSARRLPYTISSVASLQQIAPSIEEAESLGSRLNTFAKITTPMMLSGIISGAILSZ

Table 4

**ID301 (SEQ. ID. NO. 196)**

ATGAATAAGAAAAAATGATTTTAAACAAGTCTAGCCAGCGTCGCTATCTTAGGGGCTGGTTTTGTACGTCTCAGCCTACTTT  
 TGTAAGAGCAGAAAGATCTCCACAAGTTGTGCAAAAATCTTCATTAGAGAAGAAATAGAGGAAGCAAAAGCTGATA  
 CTGCCAAGAAAGATTACGAAACGGCTAAAAAGAAAGCAGAAGACGCTCAGAAAAAGTATGAAGATGATCAGAAGAACTGAG  
 GAGAAAGCTCGAAAAGAAGCAGAAGCATCTCAAAAATTGAATGATGTGGCGCTTGTGTTCAAAATGCATATAAAGAGTACCG  
 AGAAGTTCAAAATCAACGTAGTAAATATAATCTGACGCTGAATATCAGAAAAAATTAACAGAGGTCGACTCTAAAAATAGAGA  
 AGGCTAGGAAAAGACCAACAGGACTTGCAAAATAAATTTAATGAAGTAAGAGCAGTTGTAGTTCTTGAACCAAATGCGTTGGCT  
 GAGACTAAGAAAAAAGCAGAAGAAGCTAAAGCAGAAGAAAAAGTAGCTAAGAGAAAAATGATTATGCAACTCTAAAGGTAGC  
 ACTAGCGAAGAAAGAAGTAGAGGCTAAGGAACTTGAAATTGAAAAAATCAATATGAAATTTCTACTTTTGAACAAGAAGTTG  
 CTACTGCTCAACATCAAGTAGATAATTTGAAAAAACTTCTTGTGTTGCGGATCCTGATGATGGCAGAAAGTTATAGAAGCT  
 AAATTAAGAAAAAGGAGAAGCTGAGCTAAACGCTAAACAAGCTGAGTTAGCAAAAAAACAACAGAACTTGAAAAAATCTTGA  
 CAGCCTTGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTTGGATAAAAAAGCTGATGAAC  
 TTCAAAAATAAAGTTGCTGATTAGAAAAAGAAATAGTAACCTTGAAATATTACTTGGAGGGGCTGATCCTGAAGATGATACT  
 GCTGCTCTTCAAAATAAATTAGCTGCTAAAAAGCTGAGTTAGCAAAAAAACAACAGAACTTGAAAAAATCTTGGACAGCCT  
 TGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTTGGATAAAAAAGCTGATGAACCTTCAA  
 ATAAAGTTGCTGATTAGAAAAAGAAATAGTAACCTTGAAATATTACTTGGAGGGGCTGATTCTGAAGATGATACTGCTGCT  
 CTTCAAAAATAAATTAGCTTAAAAAGCTGAATTGAAAAAATCAAAAAGAATTAGATGACAGCTTAAAGTATAGGCTTGGCT  
 TGATGGAGATGAAGAAGAACTCCAGCGCCGCTCCTCAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTC  
 CAGCTCCAAAACAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTCCTCAAAACCAGAGCAACCAGCTCCTCA  
 GCTCCAAAACAGAGCAACCAGCTAAGCGGAGAAACAGCTGAAGAGCCTACTCAACCAGAAAAACAGCCACTCCAAAAAC  
 AGGCTGGAAAAACAAGAAACCGGTATGTGGTATTTCTACAATACTGATGGTTCAATGGCAATAGGTTGGCTCCAAACACCGGTT  
 CATGGTACTACCTAAACGCTAAACGGCGCTATGGCAACAGGTTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCA  
 GGTGCTATGAAGCAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATGTCAACAGCAATGGCGCTATGGCGACAGGCTG  
 GCTCCAAATACAATGGCTATGGTACTACCTCAACGCTAATGGTACTATGGCGACAGGATGGCTCCAAATACAACGGTTCTGGT  
 ATTACCTCAACGCTAATGGTATATGGCGACAGGATGGGCTAAAGTCAACGGTTCTGGTACTACCTAAACGCTAACGGTGCT  
 ATGGCTACAGGTTGGGCTAAAGTCAACGGTTCTGGTACTACCTAAACGCTAACGGTTCAATGGCAACAGGTTGGGTGAAAGA  
 TGGGATACCTGGTACTCTTGAAGCATCAGGTGCTATGAAAGCAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATG  
 TCAATGGCTTAGGTGCCCTTGCAGTCAACACAACCTGTAGATGGCTATAAAGTCAATGCCAATGGTGAATGGGTTAA

**(SEQ. ID. NO. 184)**

MNKKMILTSLASVAILGAGFVTSQPTFVRAEESPQVVEKSSLEKKYEEAKAKADTAKKDYETAKKKAEDAQKKYEDDQKRTE  
 EKARKEAEASQKLNVDVALVQNAVKEYREVQNORSKYKSDAEYQKLTTEVDSKIEKARKEQQDLQNKFNVEVRAVVVPEPNALA  
 ETKKKAEEAKAEKVKAKRYDYATLKVALAKKEVEAKELEIEKLQYEISTLEQEVATAQHVDNLKLLLAGADPDDGTEVIEA  
 KLKKGAEELNAKQAELEKQTELEKLLDSDPEGKTQDELDEKEAEAEELDKADELQNKVADLEKEISNLEILLGGADPEDDT  
 AALQNKLAAKKAELAKKQTELEKLLDSDPEGKTQDELDEKEAEAEELDKADELQNKVADLEKEISNLEILLGGADSEDDTAA  
 LQNKLATKKAELKQTELEKLLDSDPEGKTQDELDEKEAEAEELDKADELQNKVADLEKEISNLEILLGGADPEDDT  
 APKPEQPAKPEKPAEETQPEKPAETPKTGWKQENGWYFYNTDGSMAIGWLQNGSWYYLNGANGMATGWKVDGDTWYLYEAS  
 GAMKASQWFKVSDKWYVNSNGAMATGWLQYNGSWYYLNGANGMATGWLQYNGSWYYLNGANGMATGWAKVNGSWYYLNGANG  
 MATGWAKVNGSWYYLNGANGSMATGWKVDGDTWYLYEASGAMKASQWFKVSDKWYVNSNGALAVNTTVDGYKVNANGWVZ

**ID302 (SEQ. ID. NO. 197)**

ATGTTTGCATCAAAAAGCGAAAGAAAAGTACATTATTCAATTTCGTAAATTTAGTGTGGAGTAGCTAGTGTAGTTGTTGCCAG  
 TCTTGTATTGGGAAGTGTGGTTTCATGCGACAGAGAACGAGGGAGCTACCCAAGTACCCACTTCTTCTAATAGGGCAATGAAA  
 GTCCAGGCAGAACCAAGGAGAACCAACCTAAAAAATCTCGATTCAAGACGAGATAAGGCAAGGAAAGAGGTGAGGAATATGTA  
 AAAATAGTGGGTGAGAGCTATGCAAAATCACTAAAAAGCGACATACAATTACTGTAGCTCTAGTTAACGAGTTGAACAACAT  
 TAAGAACGAGTATTTGAATAAAATAGTTGAATCAACCTCAGAAAAGCCACTACAGATACTGATGATGGAGAGTCGATCAAAAG  
 TAGATGAAGCTGTGTCTAAGTTTGAAGAAGACTCATCTTCTTCGTCAAGTTTCAAGCTCTTCCACTAAACCGGAAGCTTCAGAT  
 ACAGCGAAGCCAAACAAGCCGACAGAACCCAGGAGAAAAGGTAGCAGAAGCTAAGAAGAAGGTTGAAGAAGCTGAGAAAAGAGC  
 CAAGGATCAAAAAGAAGAAGATCGTCGTAACCTACCCAACCTACTTACAAAACGCTTGAACCTGAAATTGCTGAGTCCGATG  
 TGGAAAGTTAAAAAGCGGAGCTTGAACCTAGTAAAAAGTTAAAGCTAACGAACCTCGAGACGAGCAAAAAATTAAGCAAGCAGAA  
 GCGGAAGTTGAGAGTAAACAAGCTGAGGCTACAAGGTTAAAAAAGTAAAGACAGATCGTGAAGAAGCAGAGAAGAAGCTTAA  
 ACAGAGAGCAGATGCTAAAGAGCAAGGTAAACCAAGGGGCGGGCAAAACGAGGAGTTCTTGGAGAGCTAGCAACACCTGATA  
 AAAAGAAAATGATGCGAAGTCTTCAGATTCTAGCGTAGGTGAAGAAACTCTTCAAGCCCATCCCTGAAACCGAAAAAAG  
 GTAGCAGAAGCTGAGAAGAAGGTTGAAGAAGCTAAGAAAAAAGCGGAGGATCAAAAAGAAGAAGATCGCCGTAACCTACCAAC  
 CAATACTTACAAAACGCTTGAACCTGAAATTGCTGAGTCCGATGTGGAAGTTAAAAAAGCGGAGCTTGAACCTAGTAAAGAGG  
 AAGCTAAGGAACCTCGAAACGAGGAAAAAGTTAAGCAAGCAAAAGCGGAAGTTGAGAGTAAAAAAGCTGAGGCTACAAGGTTA  
 GAAAAAATCAAGACAGATCGTAAAAAAGCAGAAGAAGCTTAAACGAAAAAGCAGCAGAAGAAGATAAAGTTAAAGAAAAACC  
 AGCTGAACAACCAACCAAGCGCGCGGCTCCAAAAGCAGAAAAACCAGCTCCAAACAGAGAAATCCAGCTGAACCAAC  
 CAAAAGCAGAAAAACAGCTGATCAACAAGCTGAAGAAGCTATGCTCGTAGATCAGAAGAAGAATATAATCGCTTGACTCAA  
 CAGCAACCGCCAAAAACTGAAAAACAGCACAACCATCTACTCCAAAAACAGGCTGGAACAAGAAAAACGGTATGTGGTACTT  
 CTACAATACTGATGGTTCAATGGCGACAGGATGGCTCCAAAAACAATGGCTCATGGTACTACCTCAACAGCAATGGCGCTATGG  
 CGACAGGATGGCTCCAAAACAATGGTTCTAGGTACTATCTAAACGCTAATGGTTCAATGGCAACAGGATGGCTCCAAAACAAT  
 GGTTCAATGGTACTACCTAAACGCTAATGGTTCAATGGCGACAGGATGGCTCAATACAATGGCTCATGGTACTACCTAAACGC  
 TAATGGTTCAATGGCGACAGGATGGCTCCAAATCAATGGTACTACCTAAACGCTAATGGTACTACCTAAACGCTAATGGTACTACCTAAACGC  
 GGCTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAAAGCAAGCCAATGGTTCAAAGTATCAGATAAA  
 TGGTACTATGTCAATGGCTCAGGTGCCCTTGCAGTCAACACAACCTGTAGATGGCTATGGAGTCAATGCCAATGGTGAATGGGT  
 AAACATA

**(SEQ. ID. NO. 185)**

MFASKSERKVVHSIRKFSVGVASVVVASLVMGSVVHATENEGATQVPTSSNRANESQAEQEQPKKLDSEDKARKEVEEYVK  
 KIVGESYAKSTKKRHTITVALVNLNNIKNEYLNKIVESTSESQQLILMMESRSKVDEAVSKFEKSSSSSSSSSSSTKPEASD  
 TAKPNKPTPEGKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLELEIAESDVEVKKAELELVKVKANEPREDEQIKQAE  
 AEVESQKAEATRLKKIKTDREEAEAEKRRADAKEQKPKGRAKRGVPELATPDKKENDAKSSDSSVGEETLPSPLKPEKK  
 VAEAEKKVEEAKKAEEDQKEEDRRNYPTNTYKLELEIAESDVEVKKAELELVKEEAEKPRNEEKVKQAEKAEVESKAEATRL  
 EKIKTDRKKAEEAEKRAAEEDKVKEPAEQPPAPAPKAEKPAPAPKPENPAEQPKAEKPADQAEEDYARRSEEEYNRLTQ



QPPKTEKPAQPSTPKTGWKQENGWMYFYNTDGSMTAGWLQNNGSWYYLNSNGAMATGWLQNNGSWYYLNSNGAMATGWLQNN  
 GSWYYLNSNGAMATGWLQYNGSWYYLNSNGAMATGWLQYNGSWYYLNSNGAMATGWVKGDTWYYLEASGAMKASQWFKVSDK  
 WYYVNGSGALAVNTTVDGYGVNANGEWVNZ

**ID303 (SEQ. ID. NO. 198)**

ATGGTAAAAAGACGTATAAGGAGAGGGACGAGAGAACCTGAAAAAGTTGTTGTTCTGAGCAATCATCTATTCTTCGTATCC  
 TGTATCTGTACATCTAACCAAGGAACAGATGTAGCAGTAGAACAGCTAAAGCAGTTGCTCCAACAACAGACTGGAAACAAG  
 AAAATGGTATGTGGTATTTTATAATACTGATGGTTCCATGGCAACAGGTTGGGTACAAGTTAATAGTTTCATGGTACTACCTC  
 AACAGCAACGGTTCTATGAAAGTCAATCAATGGTTCCAAGTTGGTGGTAAATGGTATTATGTAAATACATCGGGTGAGTTAGC  
 GGTCAATACAAGTATAGATGGCTATAGAGTCAATGATAATGGTGAATGGGTGCGTTAA

**(SEQ. ID. NO. 186)**

MVKRRIRRGTREPEKVVVPEQSSIPSYPVSVTSNQGTDVAVEPAKAVAPTTDWKQENGWMYFYNTDGSMTAGWVQVNSSWYYL  
 NSNGSMKVNQWFQVGGKYYVNTSGELAVNTSIDGYRVNDNGEWRZ

**ID304 (SEQ. ID. NO. 199)**

CTGAATACAAGTTTTGTTTCATGCTGCTGATGGGATTCAATATGTCAGAGATGATACTAGAGATAAAGAAGAGGGAATAGAGTA  
 TGATGACGCTGACAATGGGGATATTATTGTAAGTAGCGACTAAACCTAAGGTAGTAACCAAGAAAATTTCAAGTACGCGAA  
 TTCGTTATGAAAAAGATGAAACAAAAGACCGTAGTGAAAAATCCTGTTACAATTGATGGAGAGGATGGCTATGTAACCTACGACA  
 AGGACCTACGATGTTAATCCAGAGACTGGTTATGTTACCGAACAGGTTACTGTTGATAGAAAAGAAGCCAGGATACAGTTAT  
 CAAAGTTCAGCTAAAAGCAAGGTTGAAGAAGTTCTTGTTCCATTGCTACTAAATATGAAGCAGACAATGACCTTTCTGCAG  
 GACAGGAGCAAGAGATTACTCTAGGAAAGAATGGGAAAACAGTTACAACGATAACTTATAATGTAGATGGAAAGAGTGGACAA  
 GTAAGTGAAGTACTTTAAGTCAAAAAAAGACTCTCAAAACAGAGTTGTTAAAAAAGAACCAAGCCCCAAGTTCTTGTTCCA  
 AGAAATTCGAATCGAAACAGAATATCTCGATGGCCCACTCTTGATAAAAGTCAAGAAGTAGAAGAAGTAGGAGAAATTGGTA  
 AATTACTCTTACTACAATCTATACTGTAG

**(SEQ. ID. NO. 187)**

LNTSFVHAADGIQYVRDDTRDKEEGIEYDDADNGDIIVKVATKPKVVTKKISSTRIRYKDETKDRSENPTIDGEDGYVTTT  
 RTYDVPNETGYVTEQVTVDRKEATDTVIKVPKSKVEEVLVPATKYEADNDLSAGQEQEITLGNKGTVTTITYNVDSKSGQ  
 VTESTLSQKKDSQTRVVKRTRKPKQVLVQEIPIETEYLDGPTLDKSQEVEEVEIGEIKLLLLQSILZ

**ID305 (SEQ. ID. NO. 200)**

ATGAAGCTTTTGAAAAAATGATGCAAATCGCACTAGCCACATTTTTCTTCGGTTTGTAGCGACAAATACAGTATTTGCAGA  
 TGATTCTGAAGGATGGCAGTTTGTCCAAGAAAATGGTAGAACCTACTACAAAAAGGGGGATCTAAAAGAAACCTACTGGAGAG  
 TGATAGATGGGAAGTACTATTATTTTGATCCTTTATCCGGAGAGATGGTTGTGCGGTGGCAATATATACCTGCTCCACACAAG  
 GGGGTTACGATTGGTCTCTCTCCAAGAATAGAGATTGCTCTTAGACCAGATTGGTTTATTTTGGTCAAGATGGTGTATTACA  
 AGAATTTGTTGGCAAGCAAGTTTGAAGCAAAAAGCTGCTACGAATACCAACAAACATCATGGGGAAGAATATGATAGCCAAG  
 CAGAGAAACGAGTCTATTATTTGAAGATCAGCGTAGTTATCATACTTTAAAACTGGTTGGATTATGAAGAGGGTCAATTGG  
 TATTATTTACAGAAGGATGGTGGCTTTGATTGCGGCATCAACAGATTGACGGTTGGAGAGCTAGCACGTGGTTGGGTTAAGGA  
 TTACCTCTTACGTATGATGAAGAGAAGCTAAAAGCAGCTCCATGGTACTATCTAAATCCAGCAACTGGCATTATGCAACAG  
 GTTGGCAATATCTAGGTAATAGATGGTACTACCTCCATTGCTCAGGAGCTATGGCAACTGGCTGGTATAAGGAAGGCTCAACT  
 TGGTACTATCTAGATGCTGAAAATGGTATGATGAGAAGCTGGCTGGCAAAACCTTGGGAACAAATGTTACTCTCCGTTCACTC  
 AGGAGCTATGGCAACTGGTTGGTATCAGGAAAGTTGCACTTGGTACTATCTAAATGCAAGTAATGGAGATATGAAAACAGGCT  
 GTTTCCAAGTCAATGGTAACTGGTACTATGCCTATGATTAGGTGCTTTAGCTGTTAATACCACAGTAGGTGGTTACTACTTA  
 AACTATAATGGTGAATGGGTTAAGTAA

**(SEQ. ID. NO. 188)**

MKLLKMMQIALATFFGLLATNTVFADDSEGWFQVQENGRYYKKGDLKETYWRVIDGKYYYFDPLSGEMVVGWQYIIPAPHK  
 GVTIGPSPRIELRPDWFYFGQDGLVQEFVGKQVLEAKTATNTNKHGEEYDSQAEKRVYFEDQRSYHTLKTGWIYEEGHW  
 YYLQKDGGFDSRINRLTVGELARGWVKDYPLTYDEEKLKAAPWYYLNPATGIMQTGWQYLGNRWYYLHSSGAMATGWYKEGST  
 WYYLDAENGDMRTGWQNLGNKWWYYLRSSGAMATGWYQESSTWYYLNASNGDMKTGWQVNGNWWYYAYDSGALAVNTTVGGYYL  
 NYNGEWNZ

**ID306 (SEQ. ID. NO. 201)**

TTGGCTGGTAGATATGGTTCTGCTGTTCACTGTACAGAAGTGACTGCCTCAAACCTTTCAACAGTTAAAACTAAAGCTACGGT  
 TGTAAGAAAACCACTGAAAGATTTTAGAGCGTCTACGTCTGATCAGTCTGGTTGGGTGGAATCTAATGGTAAATGGTATTTCT  
 ATGAGTCTGGTGATGTGAAGACAGGTTGGGTGAAAACAGATGGTAAATGGTACTATTGAAATGACTTAGGTGTCTAGCAGACT  
 GGATTGTAAATTTCTGGTAGCTGGTATTACTTGAGCAATTGAGTGCTATGTTTACAGGCTGGGGAACAGATGGTAGCAG  
 ATGGTTCTACTTTGACGGCTCAGGAGCTATGAAGACAGGCTGGTACAAGGAAAATGGCACTTGGTATTACCTTGACGAAGCAG  
 GTATCATGAAGACAGGTTGGTTTAAAGTCGGACCACACTGGTACTATGCCTACGGTTCAGGAGCTTTGGCTGTGAGCACAA  
 ACACCAGATGGTTACCGTGTAATGGTAAATGGTGAATGGGTAAACTAG

**(SEQ. ID. NO. 189)**

LAGRYGSAVQCETVTASNLSTVKTKATVVEKPLKDFRSTSDQSGWVESNGKWYFYESGDVKTGWVKTDGKWWYLNLDLGVMT  
 GFVKFSGSWYYLNSNGAMFTGWGTDGSRWFYFDGSGAMKTGWYKENGWYLYDEAGIMKTGWFKVGPWYAYGSGALAVSTT  
 TPDGYRVNGNGEWNZ

**ID307 (SEQ. ID. NO. 202)**

ATGAAAATTTTGAAAAAATGCAAGTTGGACTGACAGTATTTTTCTTTGGTTTGCTAGGGACCAGTACAGTATTTGCAGA  
 TGATTCTGAAGGATGGCAGTTTGTCCAAGAAAACGGAAGAACCTACTACAAAAAGGGGGACCTCAAAGAAACCTACTGGCGAG  
 TGATTGATGGTAAGTACTATTATTTGATTCTCTATCTGAGAGAGATGGTTGTCGGCTGGCAATATATCCCGTTTCCATCTAAA  
 GGATGACAAATTTGGTCTTACCACAAATGGTATCAGATTGAAGAGTTTCCAAAGTCAGAGTGGTACTACTCTCGATAAAAAATGG  
 AGTGCTACAAGAGTTTGTGGTTGGAAAACATTAGAGATTAATACTAAAGACAGTGTGGGAAGAAAGTACGAGGAAAAACGTG  
 AAGATTGAGAGATAAAGAAGAGAAGCGTTATTATACGAACATTTAATCAAAATCATCTTTAGAGACAGGTTGGCTT  
 TATGATCAGTCTAACTGGTATTATCTAGCTAAGACGGAAATTAATGGAGAAAACTACCTTGGTGGTGAAAGACGTGCGGGGTG  
 GATAACAGATGATTGCTAGTTGGTACTACCTAGATCCAACAACCTGGTATTATGCAACAGGTTGGCAATATCTAGGTAATAAGT

GGTACTACCTCCGTTCTCAGGAGCAATGGCCACTGGCTGGTATCAGGAAGGTACCACTTGGTATTATTTAGACCACCCAAAT  
GGCGATATGAAAAACAGGTTGGCAAAACCTTGGGAACAAATGGTACTATCTCCGTTTCATCAGGAGCTATGGCAACTGGTTGGTA  
TCAAGATGGTTCAACTTGGTACTACCTAAATGCAGGTAATGGAGACATGAAGACAGGTTGGTTCCAGGTCAATGGCAACTGGT  
ACTATGCTTATAGCTCAGGTGCTTTGGCAGTGAATACGACCGTAGATGGCTATTCTGTCAACTATAATGGCGAATGGGTTCCG  
TAA

## (SEQ. ID. NO. 190)

MKILKKTQVGLTVFFFLGTSTVFADDSEGWQFVQENGRYYKKGLKETYWRVIDGKYYYFDSLSEMVVWGQYIPFPSK  
GSTIGPYPNIRLEGFPKSEWYYFDKNGVLQEFVWKTLEIKTKDSVGRKYGEKREDSKKEKRYTYNYFNQNSHLETGWL  
YDQSNWYYLAKTEINGENYLGERRAGWINDDSTWYYLDPTTGMQGTGWQYLGKWKYYLRSSGAMATGWYQEGTTWYYLDHPN  
ZDMKGTGWQNLGNKWKYYLRSSGAMATGWYQDGSWTWYYLNAGNGDMKTGWFWQVNGNWKYYAYSSGALAVNTTVDGYSVNYNGEWW  
Z

## ID308 (SEQ. ID. NO. 203)

ATGAAAAAGAAATTAAGTACTTGTAGCACTTGTAGGCGCTTTTTTAGGTTTGTCTATGGTATGGGAATGTTTCAAGAAAG  
TTCAGGAAATAAAATCCACTTTATCAATGTTCAAGAAGGTGGCAGTGATGCGATTATCTTGAAGCAATGGACATTTTGCCA  
TGGTGGATACAGGAGAAGATTATGATTTCCAGATGGAAGTGATTCTCGCTATCCATGGAGAGAAGGAATTGAAACGCTCTTAT  
AAGCATGTTCTAACAGACCGGTGTCTTTTCGTCGTTTGAAGGAATTTGGGTGTCCAAAACTTGATTTTATTTTGGTGACCCATAC  
CCACAGTGATCATATTGGAAATGTTGATGAATTACTGTCTACCTATCCAGTTGACCGAGTCTATCTTAAGAAATATAGTGATA  
GTCGTATTACTAATCTGAAACGTCTATGGGATAATCTGTATGGCTATGATAAGGTTTACAGAGCTGTCAGAAAAAGGTTGT  
TCAGTTATTTCAAAATATCACACAAGGGGATGCTCATTTTTCAGTTTGGGGACATGGATATTTCAGCTCTATAAATTATGAAATGA  
AACTGATTCATCGGGTGAATTAAAGAAAAATTTGGGATGACAATTTCAATTCTTGGATTAGCGTGGTGAAGTCAATGGCAAGA  
AAATTTACCTTGGGGGCGATTTAGATAATGTTTCATGGAGCAGAGACAAGTATGGTCTCTCATTGGAAGGTTGATTTGATG  
AAGTTTAAATCATCACCATGATACCAACAAATCAAATACCAAGGATTTTCAATAAAAATTTGAGTCCGAGTTTGGATTGTTCAAAC  
TTCGGATAGTCTACCTTGGAAAAATGGTGTGATAGTGAGTATGTTAATTTGGCTCAAAGAACGAGGAATTGAGAGAATCAACG  
CAGCCAGCAAAGACTATGATGCAACAGTTTTTGTATTTGAAAAAGACGGTTTTGTCAATATTTCAACATCTTCAAGCCGATT  
CCAAGTTTTCAGAGCTGGTTGGCATAAGAGTGATATGGGAACCTGGGTGATCAAGCGCTGATTCTACAGGAGAGTATGCTGT  
CGGTTGGAATGAATCGAAGGTGAATGGTATTACTTTAACCAAAACGGGTATCTTGTACAGAATCAATGGAAAAATGGAACA  
ATCATTTGTTCTATTTGACAGACTCTGGTGTCTCTGCTAAAAATTTGAAGAAAAATCGCTGGAATCTGGTATTATTTTAAACAA  
GAAACCCAGATGGAATTTGGTGGATTCAAGATAAAGAGCAGTGGTATTATTTGGATGTTGATGGTTCTATGAAGACAGGATG  
GCTTCAATATATGGGGCAATGGTATTACTTTGCTCCATCAGGGGAAATGAAATGGGCTGGGTAAGATAAAGAAAACTGGT  
ACTATATGGATTCTACTGGTGTCTATGAAGACAGGTGAGATAGAAGTTGCTGGTCAACATTATTATCTGGAGACTCAGGAGCT  
ATGAAGCAAGGCTGGCATAAAAAGGCAAATGATTGGTATTTCTACAAGACAGACGGTTACAGAGCTGTGGGTTGGATCAAGGA  
CAAGGATAAATGGTACTTCTTGAAGAAAAATGGTCAATTACTTTGTAACGGTAAGACACCAGAAGGTTTACTGTGGATTCAA  
GTGGTGCCTGGTTAGTGATGTTTCGATCGAGAAATCTGCTACAATTAAGCTACAAGTCAATGAGAAATGAAAGAAATCCAAA  
GAAGTAGTGAAGAAAGGATCTTGAAGATAAAGAAACGAGTCAACATGAAAGTGTACAAATTTTTCACTAGTCAAGATTTGAC  
ATCCTCAACTTCACAAAGCTCTGAAACGAGTGTAACAAATCGGAATCAGAACAGTAG

## (SEQ. ID. NO. 191)

MKKKLTSLALVGAFLGLSWYGNVQAQESSGNKIHFINVQEGGSDAIILESNGHFAMVDTGEDYDFPDGSDSRYPWREGIETSY  
KHVLTDRVFRRLKELGVQKLDLILVTHTHSDHIGNVDELLSTYPVDRVYLKKYSRSRTNSERLWDNLGYDKVLQTAEEKGV  
SVIQNITQGDHAFQFGMDIQLYNYENETDSSGELKKIWDNNSNLSISVVKVNGKKIYLGDDLNDNVHGEDYKGPLIGKVDLM  
KFNHHDHNTKSNKDFIKNLSPLIVQTSLSLPWKNGVDSEYVNWLERGIERINASKDYDATVFDIRKDGFTVNIISTSYKPI  
PSFQAGWHKSYAGNWWYQAPDSTGEYAVGWNEIEGEWYFNFQGTILLQNWKKWNNHWFYLTDSGASAKNWKKIAGIWIYFNF  
ENQMEIGWIQDKQWYLLDVGSMKTGWLYMGQWYFAPSGEMKMGWVKDKETWYMDSTGVMKTGEIEVAGQHYYLEDGSA  
MKQGWHKKANDWYFYKTDGSRVAGWIKDKDKWYFLKENGQLLVNGKTPGYTVDSGAWLVDVSIKSAITIKTTSHEIKESK  
EVVKKDLNENKETSQHSVTFNSTSQDLTSSTSSQSSSVNKSESEQZ

## ID309 (SEQ. ID. NO. 204)

ATGGAAATTAATGTGAGTAAATTAAGAACAGATTTGCCTCAAGTCGGCGTGCAACCATATAGGCAAGTACACGCACACTCAAC  
TGGGAATCCGCAATGAGCGTACAGAATGAAGCGGATATCACTGGCGGAAGACCCAGAATTAGGTTTTTCTCGCACATTG  
TTGGGAACGGTTGCATCATGCAGGTAGGACCTGTTGATAATGTTGCTGGCGACGTTGGGGGCGGTTGGAATGCTGAGACCTAT  
GCAGCGGTTGAATGATTGAAAGCCATTCAACCAAAGAAGAGTTTCATGACGGAATACCGCCTTTATATCGAATCTTACGCAA  
TCTAGCAGATGAAGCAGGTTTGGCGAAACGCTTGATACAGGAGTTTAGCTGGAATTAACCGCACGAGTATTGCACGAATA  
ACCAACCAACCAACCACTCAGACACCGTTGACCTTTATCCATATCTTGTCTAAATGGGGCATTAGCCGTGAGCAGTTTAAAGCAT  
GATATTGAGAACGGCTTGACGATTGAACAGGCTGGCAGAAGATGACACTGGCTACTGGTACGTACATTAGACCGGCTCTTA  
TCCAAAAGACAAGTTTGAAGAAATCAATGGCACTTGGTACTACTTTGACAGTTCAAGGCTATATGCTTGCAGACCGCTGGAGGA  
AGCACACAGACGGCAACTGGTACTGGTTGACAACTCAGGCGEMKMGWVKDKETWYMDSTGVMKTGEIEVAGQHYYLEDGSA  
TATTTCAACGAAGAAGGTGCCATGAAGACAGGCTGGGTCAAGTACAAGGACACTTGGTACTACTTAGACGCTAAAGAAGGCGC  
CATGGTATCAAATGCCTTTATCCAGTCAGCGGACGGAACAGGCTGGTACTACTCAAACAGACGGAACACTGGCAGACAAGC  
CAGAATTCACAGTAGAGCCAGATGGCTTGATTACAGTAAATAA

## (SEQ. ID. NO. 192)

MEINVSKLRDTLPQVGVPYRQVHAHSTGNPHSTVQNEADYHWRKDPGLGFFSHIVGNGCIMQVGPVDNGAWDVGGGWNAETY  
AAVELIESHSTKEEFMTDYRLYIELLRNLADEAGLPKTLDTGSLAGIKTHEYCTNNQPNHSDHVDPPYLAKEWISREQFKH  
DIENGLTIETGWQKNDTGYYVHSDGSYPKDKFEKINGTWYFDSSGYMLADRWKHTDGNWYWFNDNGEMATGWKKIADKWY  
YFNEEGAMKTGWVKYKDTWYYLDAKEGAMVSNAFIQSADGTGWYYLKPDPGLADKPEFTVEPDGLITVKZ

## ID310 (SEQ. ID. NO. 205)

ATGGGCACAACAGGATTTACAATAATTGACTTAATTATCTTGATTGTTTATTTACTTGGCGGTGTTGGTTGCAGGTATCTATTT  
CTCTAAAAAGAGATGAAAGGAAAGAGTTCTTTAAAGGAGATGGTTCCGTTCCCTTGGTATGTTACTTCCGTTATCCATTTTGG  
CCACAATGCTCAGTCCGATTTCTTTGGGACTCGCTGGTACTCTTATCAGGTAGCTGGATTATGAGTTGCTGAGTTGCTCAATTA  
GGGATGGTAGTAGCTATTCCACTGACAATTCGTTTTATCTTACCTATCTTTGCACGGATAGACATCGATACGGCATATGATTA  
CTTGGATAAACGTTTAAATCTAAAGCACTTCGTATTTTTCAGCACTCTTGTATTATTTATCAATTTGGGACGTTATGCTA  
TCATTATGTACCTCCCATCAGCTGGTTATCAGTATTGACAGGAATGGCTCAATATTTTGGATTATTTTGGTGGTGGTGGT  
GCAATTGTTTATTTCTTACTGTTGGTCTAAATCCGTATTATGGACAGACTTTATTCAAGGTGTGATTCTGATTAGTGGTGT

CGTTTTAGCTTTATTTGTACTGATTGCTAATATTAAGGTGGCTTTGGTGCAGTAGCAGAAACATTAGCAAACGGGAAATTCC  
 TTGCTGCAAATGAAAACTTTTCGATCCTAAGTCTTCTTCAAACTCCATCTTTTTAATTGTGATGGGTTCCAGGCTTTACAATC  
 TTGTCTTCTCTATGCTTCATCTCAAGATTTGGTTCACGTTTTACTACAACACAAATATTAAGAACTTAATAAGATGTTGTT  
 CACAAACGGTGTGTTTGTCACTTGCAACTGCAACAGCTCTTTTACTTGATTGGTACAGGCTGTACGTATTCTATCAAGTACAAA  
 ATGCAGATAGTGCAGCTAGCAATATCCCTCAAGACCAATCTTTATGCTATTTATGCAATACCAGTTACCAGTGATCACA  
 GGTGTTGATCTTGGCAGCGATTATGCAGCATCTCAATCAACTATTTCAACAGGTTTGAACCTCTGTTGCAACTTCATGGACATT  
 GGATATTCAGATGTCTATTCTAAAAATATGTGAGACAATCGTCGTACGAAAATTGCACAATTCGTATCTCTAGCAGTAGGTT  
 TATTCTCAATTGGTGTGTTTCCATTGTCATGGCTCACTCAGATATTAATCTGCATACGAATGGTTCAATAGTTTCATGGGACTT  
 GTACTTGGTCTACTTGGTGGTGTATTTATCTTGGATTGTGTTCTAAAAAAGCAAATAAACAAGGTGCTTATGCAGCGCTGAT  
 TGTATCAACCATCGTCATGGTATTTATTAATACTTCTCTCCCAACAGCTGTTAGCTACTGGGCATATTCTATTGATTTCAA  
 TCTCTGTATCAGATAGTTTCAGGTTATATTGTATCTGTTCTTACTGGAAATAAAGTATCTGCACCTAAATATACAACGATTCTAT  
 GATATTACAGAAATTAAGCGGATTCAAGTTGGGAAGTTTCGTCACCTAA

## (SEQ. ID. NO. 193)

MGTTGFTIIDLIIILIVYLLAVLVAGIYFSKKEMKGKEFFKGDGVSVPWYVTSVSIFATMLSPISFLGLAGSSYAGSWILWFAQL  
 GMVVAIPLTIRFILPIFARIDIDTAYDYLDKRFNSKALRIISALLFIYQLGRMSIMYLPSAGLSVLTGIDINILIIILMGVV  
 AIVYSYTGGLKSVLWTDPIQGVILISGVVLLALFVLIANIKGGFGAVAETLANGKFLAANEKLFDPNLLSNSIFLIVMSGFTI  
 LSSYASSQDLVQRFRTTQNIKKLNKMLFTNGVLSLATATVFYILGTGLYVYFQVQNADSAASNIPODQIFMYFIAYQLPVGIT  
 GLILAAIYAASQSTISTGLNSVATSWTLDIQDVISKNMSDNRRTKIAQFVSLAVGLFISGVSVMAHSDIKSAYEWFNSFMGL  
 VLGLLGGVFIILGFVSKKANKQGAYAALIVSTIVMVFIKYFLPPTAVSYWAYSLISISVSVVSGYIVSVLTGNKVSAPKYTTIH  
 DITEIKADSSWEVRHZ

## ID311 (SEQ. ID. NO. 206)

ATGAAAAATTAATAAAAAATATCTAGCAGGTTCAAGTGGCAGTCCCTGCCCCTAAGTGTGTTGTTCTCTATGAGCTTGGTGCCTACCA  
 AGCTGGTCAGGATAAGAAAGAGTCTAATCGAGTTGCTTATATAGATGGTGTATCAGGCTGGTCAAAGGCGAGAAAACTTGACAC  
 CAGATGAAGTCAGTAAGAGGGAGGGATCAACGCCGAAACAAATCGTCATCAAGATTACGGATCAAGGTTATGTGACCTCTCAT  
 GGAGACCATTATCATTACTATAATGGCAAGGTCCCTTATGATGCATCATCAGTGAAGAGCTCCTCATGAAAGATCCGAATTA  
 TCAGTTGAAGGATTGACAGATTGTCAATGAAATCAAGGGTGGTTATGTCTATCAAGGTAGACGGAAAACTACTATGTTTACCTTA  
 AGGATGCAGCTCATGCCGATAATATTCGGACAAAAGAGATTAACAGCTCAGAAGCAGGAACGAGTCAATCAACGGGTCA  
 GGAGCTAACGATCATGCAGTAGCTGCAGCCAGAGCCCAAGGACGCTATACAACGGATGATGGGTATATCTTCAATGCATCTGA  
 TATCATTGAGGACACGGGTGATGCTTATATCGTTCTCCTCAGCGGACCATTAACATTACATTCTCTAAGAATGAGTTATCAGCTA  
 GCGAGTTAGCTGCTGCAGAAGCCTATTGGAATGGGAAGCAGGGAATCTCGTCTTCTTCAAGTTCTAGTTTATAATGCCAAATCCA  
 GCTCAACCAAGATTGTGAGAGAACCACAACTCTGACTGTCTCACTCAACTTATCATCAAAATCAAGGGGAAAACTTTCAAGCCT  
 TTTACGTGAATTGTATGCTAAACCCCTTATCAGAAGCCCATGTGGAATCTGATGGCCTTATTTTCGACCCAGCGCAAATCACAA  
 GTCGAACCCGACAGAGGTGATGCTGCTCCCTCATGGTAACCATTAACCACTTATCCCTTATGAACAAATGTCTGAATTGGAAAAA  
 CGAATTGCTCGTATTTATTTCCCTTCGTTATCGTTCAAACCATTTGGGTACCAGATTCAAGACCAGAACAACCAAGTCCACAATC  
 GACTCCGGAACCTAGTCCAAGTCCGCAACCTGCACCAATCCTCAACCAGCTCCAAGCAATCCAATTGATGAGAAATTGGTCA  
 AAGAAGCTGTTGCAAAAGTAGGCGATGGTTATGTCTTTGAGGAGAATGGAGTTTCTCGTTATATCCCAGCCAAAGGATCTTTCA  
 GCAGAAACAGCAGCAGGCGATTGATAGCAAACTGGCCAAAGCAGGAAAGTTTATCTCATAAGCTAGGAGCTAAGAAAAGTACCT  
 CCCATCTAGTGATCGAGAATTTTACAATAAGGCTTATGACTTACTAGCAAGAATTCACCAAGATTTACTTGATAATAAAGGTC  
 GACAAGTTGATTTTGGAGCTTTGGATAACCTGTTGGAACGACTCAAGGATGTCCCAAGTGATAAAGTCAAGTTAGTGGATGAT  
 ATTCTTGCCTTCTTAGCTCCGATTCTGTCATCCGAACGTTTAGGAAAACCAATGCGCAAATTACCTACACTGATGATGAGAT  
 TCAAGTAGCCAAGTTGGCAGGCAAGTACACAACAGAAGACGGTTATATCTTTGATCCTCGTGATATAACCAAGTATGAGGGG  
 ATGCCTATGTAACCTCACATATGACCCATAGCCATAGCAGATTAAGAAAGATAGTTTGTCTGAAGCTGAGAGAGCGGCAGCCAG  
 GCTTATGCTAAAGAGAAAGGTTGACCCCTCCTTCGACAGACCATCAGGATTCAGGAAATCTGAGGCAAAAGGAGCAGAAAGC  
 TATCTACAACCGCTGAGGAGCAGTAAAGAGGCTGACCTTATGATCGTATGCTTACAACTTCAATATCTGATAGAGTCAAAA  
 ACGGTAGTTTAAATCATACCTCATTATGACCATTACCATAACATCAAATTTGAGTGGTTTGACGAAGGCCTTTATGAGGCACCT  
 AAGGGGTACTCTTGGAGATCTTTTGGCGACTGTCAAGTACTATGTGCAACATCCAACGAAACGCTCCGCTATCAGATAATGG  
 TTTGGTAACCGTAGCGACCATGTTCAAAGAAACAAATGTTCAAGCTGATACCAATCAAACGGAACCAAGCAGGAGGAGA  
 AACCTCAGACAGAAAAACCTGAGGAAGAAACCCCTCGAGAAGAGAAACCGCAAAGCAGAGAAACAGAGTCTCCAAAACCAACA  
 GAGGAACCAAGAAATCACCAGAGGAATCAGAAGAACCTCAGGTGAGAGCTGAAAAGGTTGAAGAAAACTGAGAGAGGCTGA  
 AGATTACTTGGAAAAATCCAGGATCCAATTCAAGTCCAATGCCAAGAGACTCTCACAGGATTAATAAATTTACTAT  
 TTGGCACCCAGGACAACAATACTATTATGAGCAGAAGCTGAAAAACTATTGGCTTTATTAAGGAGAGTAAGTAA

## (SEQ. ID. NO. 194)

MKINKKYLGSVAVLALSVCSYELGRHQAGQDKKESNRVAYIDGDAQQKAENLTPDEVSKREGINAQIVIKITDQGYVTS  
 GDHYHYNGKVPYDAIISEELLMKDPNYQLKDSIDVNEIKGGYVIKVDGKYVYVYLKDAHADNIRTKKEIKRQKQERSHNHGS  
 GANDHAVAARAQGRYTTDDGYIFNASDIIEDTGDYIVPHGDHYHYIIPKNELSASELAAEA YWNGKQGRSPSSSSSYNANP  
 AQPRLSENHNLVTPTYHQNGENISSLLRELYAKPLSERHVESDGLIFDPAQITSRARGVAVPHGNHYHFIPEQMSLEK  
 RIARIIPLYRNSNHWPDSRPEQSPQSTPEPSPSPQAPNPQAPSNPIDKLVKEAVRKVGDDYVFEENGVSRYIPAKDLS  
 AETAAGIDS KLAKQESLSHLKLGAKKTDLPSSDREFYNKAYDLLARIHQDLDLNKGRQVDFEALDNLRLKDVPSDKVLVDD  
 ILAFLAPIRHPERLGPKNQAITYTDBEIQVAKLAGKYTTEDGYIFDPRDITSDEGDAYVTPHMTSHHWIKKDSLSEABRAAAQ  
 AYAKEKGLTPPSTDHQDSGNTAEKGAELIYNRVKAAKVP LDRMPYNLYQTVFVKNGLSIIIPHYDHYNIKFEFWFDEGLYEAP  
 KGYTLEDLLATVKYVVEHPNERPHSDNGFGNASDHVQRNKGQADTNQTEKPEEKPTTEKPEETPREKPPQSEKPEPKPT  
 EEPESPESEEPQVETEKVEEKLREAEDLLGKIQDPPIKSKNAKETLTGLKNLLFGTQDNNTIMAEAEKLLALLKESKZ

## ID312 (SEQ. ID. NO. 207)

ATGAGGGGATTGGTTAGAGTGCATTTATTGCCTGTATTGGCGATTACAAGCTATCTAACTTACTACGCCTATTCTTCAACA  
 GCAAGTAAACAAATGGGCTGACAAAGGCAAAATAAGCGGAAAAAGGGGCAATTGCTAACTACTCTTGTCCATAACATGAATA  
 AGCGTATTTTGAATATGGCGTAGCTATCCAGGTAATAACAACAACCGCTAATGATGTCTATGCTCCACGCAACAGCAA  
 AAAGAAAAGGCTGCTGTCAAATACTTAGACAACAAGAATAAAACAGTTTCTTGATTATTTAGATGCTCTGGATCAATCAAA  
 TTATGAGAATCTATTGATGTTGTTCTGTATAAGACTTTATGGCCACTGGTTGGCGGTATTAGTGAGGCTCTGGCTCTTGAAT  
 GGTCTGATTAATGACTAGAAAGCGGTGTTATCAGCATCAATAAGACACTTAAACCGCTATCAGGAAATAAATCAAGTAAATCA  
 AGCGCTGGTTATCGTGATATACCAATAGACAAAGCCACATTACTTTTACTGAAACAATACAAAAACCGTCAACAAATTCAGTC  
 TTGGAATTAGGCCGATCTGAAACAGTTGTATTCTCTGTATTTACGGAGAAATATGCTTATGCTTGTAACCTACGCAACGCC  
 TAAATAAGCATTTTGATGCTGCTGGAGTAACTAACGTATCATTTCTGTTTCCGCCATACATACTACTATGATGCTCTAT

GCTCAGGTTAGCCCGAAAGATGTTTCAGTATAGATTAGGCCACTCTAATTTAATGATCACTGAAAATACTTACTGGCATACTAA  
CCAAGAGAATGCAAAAAAGCCGTCTCAAATTATGAAACAGCTATCAACAATTTATAA

(SEQ. ID. NO. 195)

MEGLVRVHLLPVFGDYKLSKLTTPILQQQVNWADKANKGEGAFANYSLHNMNKRILKYGVAIQVIQYNPANDVIVPRKQQ  
KEKAAVKYLDNKELKQFLDYLDALDQSNYENLFDVVLYKTLLATGCRISEALALEWSDIDLESGVISINKTLNRYQEINSPKS  
SAGYRDIPIDKATLLLLLKQYKNRQQIQSWKLGRSETVVFSVFTEKYAYACNLRLKRLNKHFDAAGVTNVSFHGFRTHTTMMMLY  
AQVSPKDVQYRLGHSNLMITENTYWHNTQENAKKAVSNYETAINNLLZ

At Page 1, 2<sup>nd</sup> full paragraph:

### **Background of the Invention**

*Streptococcus pneumoniae*, commonly referred to as the pneumococcus, is an important pathogenic organism. The continuing significance of *Streptococcus pneumoniae* infections in relation to human disease in developing and developed countries has been authoritatively reviewed (Fiber, G.R., Science, **265**:1385-1387 (1994)). That indicates that on a global scale this organism is believed to be the most common bacterial cause of acute respiratory infections, and is estimated to result in 1 million childhood deaths each year, mostly in developing countries (Stansfield, S.K., *Pediatr. Infect. Dis.*, **6**:622 (1987)). In the USA it has been suggested (Breiman *et al*, *Arch. Intern. Med.*, **150**:1401 (1990)) that the pneumococcus is still the most common cause of bacterial pneumoniae, and that disease rates are particularly high in young children, in the elderly, and in patients with predisposing conditions such as asplenia, heart, lung, and kidney disease, diabetes, alcoholism, or with immunosuppressive disorders, especially AIDS. These groups are at higher risk of pneumococcal septicaemia and hence meningitis and therefore have a greater risk of dying from pneumococcal infection. The pneumococcus is also the leading cause of otitis media and sinusitis, which remain prevalent infections in children in developed countries, and which incur substantial costs.

At Page 3, 2<sup>nd</sup> full paragraph:

### **Brief Summary of the Invention**

Thus, in a first aspect the present invention provides a *Streptococcus pneumoniae* protein or polypeptide having a sequence selected from those shown in Table 2 herein.

At Page 3, 2<sup>nd</sup> full paragraph:

**Detailed Description of the Invention**

As discussed herein, the proteins and polypeptides of the invention are useful as antigenic material. Such material can be “antigenic” and/or “immunogenic”. Generally, “antigenic” is taken to mean that the protein or polypeptide is capable of being used to raise antibodies or indeed is capable of inducing an antibody response in a subject. “Immunogenic” is taken to mean that the protein or polypeptide is capable of eliciting a protective immune response in a subject. Thus, in the latter case, the protein or polypeptide may be capable of not only generating an antibody response and in addition non-antibody based immune responses.

At page 101, line 1:

~~CLAIMS~~ What is claimed is: